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ISSN 2292-0838 (en ligne)
Robust Dynamic Panel Data Models Using $\varepsilon$-contamination *

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Abstract/Résumé

This paper extends the work of Baltagi et al. (2018) to the popular dynamic panel data model. We investigate the robustness of Bayesian panel data models to possible misspecification of the prior distribution. The proposed robust Bayesian approach departs from the standard Bayesian framework in two ways. First, we consider the $\varepsilon$-contamination class of prior distributions for the model parameters as well as for the individual effects. Second, both the base elicited priors and the $\varepsilon$-contamination priors use Zellner (1986)'s $g$-priors for the variance-covariance matrices. We propose a general "toolbox" for a wide range of specifications which includes the dynamic panel model with random effects, with cross-correlated effects à la Chamberlain, for the Hausman-Taylor world and for dynamic panel data models with homogeneous/heterogeneous slopes and cross-sectional dependence. Using a Monte Carlo simulation study, we compare the finite sample properties of our proposed estimator to those of standard classical estimators. The paper contributes to the dynamic panel data literature by proposing a general robust Bayesian framework which encompasses the conventional frequentist specifications and their associated estimation methods as special cases.

Keywords/Mots-clés: Dynamic Model, $\varepsilon$-contamination, $g$-priors, Type-II Maximum Likelihood Posterior Density, Panel Data, Robust Bayesian Estimator, Two-Stage Hierarchy

JEL Codes/Codes JEL: C11, C23, C26

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* This paper is written in honor of M. Hashem Pesaran for his many contributions to econometrics. In particular, heterogeneous panel data, Bayesian estimation of dynamic panel data models, random coefficient models for panel data and cross-section dependence in panels. We would like to thank Jean-Michel Etienne for help and support with Stata Mata codes. The usual disclaimers apply.

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1. Introduction

The dynamic panel data model allows for feedback from lagged endogenous values and have been used in many empirical studies. The most popular estimation method is the generalized method of moments (GMM) with many variants, the best known being the Arellano-Bond difference GMM (Arellano and Bond (1991)) and the Blundell-Bond system GMM (Blundell and Bond (1998)) (see the surveys by Harris et al. (2008) and Bun and Sarafidis (2015) to mention a few). Despite its optimal asymptotic properties, Bun and Sarafidis (2015) and Moon et al. (2015), among others, argue that the finite sample behavior of the GMM estimator can be poor due to weakness and/or abundance of moment conditions and dependence on crucial nuisance parameters. Several alternative inference methods derived from inconsistent least squares (LS) or likelihood based procedures have been proposed. These include modifications of the profile likelihood (Lancaster (2002), Dhaene and Jochmans (2011, 2016)) or estimation methods based on the likelihood function of the first differences (Hsiao et al. (2002), Binder et al. (2005), Hayakawa and Pesaran (2015)).

While GMM estimation is very attractive because of its flexibility, other promising methods remain underrepresented in empirical work. Examples are bias-correction procedures for the fixed-effects dynamic panel estimator proposed by Kiviet (1995), Bun (2003), Bun and Kiviet (2003), Everaert and Pozzi (2007), and Everaert (2013) among others. Estimation of dynamic panel data models with heterogeneous slopes and/or cross-sectional dependence has also been investigated by Chudik and Pesaran (2015a,b), using the common correlated effects (CCE) approach of Pesaran (2006), and by Moon and Weidner (2015, 2017), who studied linear models with interactive fixed effects.

Quasi-maximum likelihood (QML) methods have been also proposed to circumvent this bias by modeling the unconditional likelihood function instead of conditioning on the initial observations. While this requires additional assumptions on the marginal distribution of the initial observations, the QML estimators are an attractive alternative to other estimation approaches in terms of efficiency and finite-sample performance if all the assumptions are satisfied. QML estimators can be characterized as limited-information maximum likelihood estimators that are special cases of a structural equation modeling or full information maximum-likelihood approach with many cross-equation restrictions. For dynamic models with random effects, we must be explicit about the non-zero correlation between the individual-specific effects and the initial conditions (see Anderson and Hsiao (1982), Bhargava and Sargan (1983), Alvarez and Arellano (2003), Hsiao and Pesaran (2008), Moral-Benito (2012, 2013), Kripfganz (2016), Bun et al. (2017), Moral-Benito et al. (2019)).

The widely used difference GMM estimator suffers from finite sample bias when the number of cross-section observations is small. Moreover, some have expressed concern in recent years that many instrumental variables of the type considered in panel GMM estimators such as Arellano and Bond (1991) may be invalid, weak or both (see for instance Bazzi and Clemens (2013) and Kraay (2015)). Based on the same identifying assumption, some alternatives have been proposed in the literature (e.g. Ahn and Schmidt (1995), Hansen et al. (1996), Hsiao et al. (2002), Moral-

\footnote{For the dynamic fixed-effects model, see for instance Hsiao et al. (2002).}

\footnote{In a Gaussian dynamic linear mixed model: \( y_{it} = \rho y_{i,t-1} + X_{it}' \beta + W_{it}' b_i + u_{it}, i = 1, ..., N, t = 2, ..., T \), as in our case (see eq.(8) in section (2)), maximum likelihood analysis is subject to an initial condition problem if the permanent subject effects \( b_i \) and the initial observations are correlated. In case of such correlation, possible options are a joint random prior (e.g., bivariate normal) involving \( b_i \) and the first disturbance \( u_{i1} \) (Dorsett (1999)), or a prior for \( b_i \) that is conditional on \( y_{i1} \), such as \( b_i \mid y_{i1} \sim N(\varphi y_{i1}, \sigma_b^2) \) (see Hirano (2002), Wooldridge (2005)) and Congdon (2010)).}
Benito (2013), to mention a few). Maximum likelihood estimators, asymptotically equivalent to the Arellano and Bond (1991) estimator, have recently been proposed and are strongly preferred in terms of finite sample performance (Moral-Benito et al. (2019)).

Bayesian analysis for dynamic panel data models have also been proposed (see for instance Hsiao et al. (1999), Lancaster (2002), Hsiao and Pesaran (2008), Koop et al. (2008), Juárez and Steel (2010), Tsai (2016), Liu et al. (2017), Liu et al. (2018), Bretó et al. (2019), Pacifico (2019)). Some consider that the process which generates the initial observation $y_{i0}$ of the dependent variable for each individual $i$ has started a long time ago (e.g., Juárez and Steel (2010). Others derive the estimators under the assumption that $y_{i0}$ are fixed constants (e.g., Hsiao et al. (1999), Hsiao and Pesaran (2008)). Yet others consider that the initial value is generated from the finite past using state space forms (e.g., Liu et al. (2017)), or use the Prais-Winsten transformation for the initial period. A simplifying approach, more feasible for large $T$, is to condition on the first observation in a model involving a first-order lag in $y$, so that $y_{i1}$ is nonstochastic (Hjellvik and Tjøstheim (1999), Bauwens et al. (2005)). Geweke and Keane (2000) and Lancaster (2002) consider Bayesian approaches to the dynamic linear panel model in which the model for period 1 is not necessarily linked to those for subsequent periods in a way consistent with stationarity (see also Congdon (2010)).

This brief overview seems to confirm the strong comeback of ML methods and associated Bayesian approaches for dynamic panel data models. MCMC holds some advantages over ML or QML estimation. Su and Yang (2015) and Yu et al. (2008) have discussed issues involved in maximizing a concentrated version of the likelihood function that could involve trivariate optimization over the parameters and subject to stationarity restrictions. This type of constrained optimization may lead to local optima and may produce misleading inference. In our earlier paper (Baltagi et al., 2018), which considered a static panel data model, we argued that the Bayesian approach rests upon hypothesized prior distributions (and possibly on their hyperparameters). The choice of specific distributions is often made out of convenience. Yet, it is well-known that the estimators can be sensitive to misspecification of the latter. Fortunately, this difficulty can be partly circumvented by use of the robust Bayesian approach which relies upon a class of prior distributions and selects an appropriate one in a data dependent fashion. This paper extends our earlier paper to the popular dynamic panel data model and studies the robustness of Bayesian panel data models to possible misspecification of the prior distribution in the spirit of the works of Good (1965), Dempster (1977), Rubin (1977), Hill (1980), Berger (1985), Berger and Berliner (1984) and Berger and Berliner (1986). In particular, it is concerned with the posterior robustness which is different from the robustness à la White (1980). The objective of our paper is to propose a robust Bayesian approach for dynamic panel data models which departs from the standard Bayesian one in two ways. First, we consider the $\varepsilon$-contamination class of prior distributions for the model parameters (and for the individual effects). Second, both the base elicited priors and the $\varepsilon$-contamination priors use Zellner (1986)’s $g$-priors rather than the standard Wishart distributions for the variance-covariance matrices. We propose a general “toolbox” for a wide range of specifications such as the dynamic panel model with random effects, or with cross-correlated effects à la Mundlak or à la Chamberlain, for the Hausman-Taylor world or for dynamic panel data models with homogeneous/heterogeneous slopes and cross-sectional dependence. The paper contributes to the dynamic panel data literature by proposing a general robust Bayesian framework which encompasses all the above-mentioned conventional frequentist specifications and their associated estimation methods as special cases.

Section 2 gives the general framework of a robust linear dynamic panel data model using $\varepsilon$-contamination and derives the Type-II maximum likelihood posterior mean and the variance-
covariance matrix of the coefficients in a two-stage hierarchy model. Section 3 investigates the finite sample performance of our robust Bayesian estimator through extensive Monte Carlo experiments. The simulation results underscore the relatively good performance of the two-stage hierarchy estimator as compared to the standard frequentist estimation methods. Section 4 gives our conclusion.

2. A robust linear dynamic panel data model

2.1. The static framework

Baltagi et al. (2018) considered the following Gaussian static linear mixed model:

\[ y_{it} = X'_{it} \beta + W'_{it} b_i + u_{it}, \quad i = 1, ..., N, \quad t = 1, ..., T, \tag{1} \]

where \( X'_{it} \) is a \((1 \times K_x)\) vector of explanatory variables including the intercept, and \( \beta \) is a \((K_x \times 1)\) vector of parameters. \( t \) is the faster index (primal pooling). Furthermore, let \( W'_{it} \) denote a \((1 \times K_2)\) vector of covariates and \( b_i \) a \((K_2 \times 1)\) vector of parameters. The subscript \( i \) of \( b_i \) indicates that the model allows for heterogeneity on the \( W \) variables. The distribution of \( u_{it} \) is parametrized in terms of its precision \( \tau \) rather than its variance \( \sigma^2_u \) (\( = 1/\tau \)).

Following the seminal papers of Lindley and Smith (1972) and Smith (1973), various authors including Chib and Carlin (1999), Koop (2003), Chib (2008), Greenberg (2008), Zheng et al. (2008), and Rendon (2013) have proposed a very general three-stage hierarchy framework

First stage:

\[ y = X \beta + Wb + u, \quad u \sim N(0, \Sigma), \Sigma = \tau^{-1} I_{NT} \]

Second stage:

\[ \beta \sim N(\beta_0, \Lambda_\beta) \quad \text{and} \quad b \sim N(b_0, \Lambda_b) \tag{2} \]

Third stage:

\[ \Lambda_b^{-1} \sim \text{Wish}(v_b, R_b) \quad \text{and} \quad \tau \sim G(\cdot), \]

where \( y = (y_{1,1}, ..., y_{1,T}, ..., y_{N,1}, ..., y_{N,T})' \) is \((NT \times 1)\). \( X \) is \((NT \times K_x)\), \( W \) is \((NT \times K_2)\), \( u \) is \((NT \times 1)\) and \( I_{NT} \) is a \((NT \times NT)\) identity matrix. The parameters depend upon hyperparameters which themselves follow random distributions. The second stage (also called fixed effects model in the Bayesian literature) updates the distribution of the parameters. The third stage (also called random effects model in the Bayesian literature) updates the distribution of the hyperparameters. The random effects model simply updates the distribution of the hyperparameters. The precision \( \tau \) is assumed to follow a Gamma distribution and \( \Lambda_b^{-1} \) is assumed to follow a Wishart distribution with \( v_b \) degrees of freedom and a hyperparameter matrix \( R_b \) which is generally chosen close to an identity matrix. In that case, the hyperparameters only concern the variance-covariance matrix of the \( b \) coefficients and the precision \( \tau \). As is well-known, Bayesian methods are sensitive to misspecification of the distributions of the priors. Conventional proper priors in the normal linear model have been based on the conjugate Normal-Gamma family because they allow closed form calculations of all marginal likelihoods. Likewise, rather than specifying a Wishart distribution for the variance-covariance matrices as is customary, Zellner’s \( g \)-prior \((\Lambda_\beta = (\tau g X'X)^{-1} \text{ for } \beta \text{ or } \Lambda_b = (\tau h W'W)^{-1} \text{ for } b) \) has been widely adopted because of its computational efficiency in evaluating marginal likelihoods and because of its simple interpretation arising from the design matrix of observables in the sample. Since the calculation of marginal likelihoods using a mixture of \( g \)-priors involves only a one-dimensional integral, this approach provides an attractive computational solution that made the original \( g \)-priors popular while insuring robustness to misspecification of \( g \) (see Zellner (1986) and Fernández et al. (2001)).
To guard against mispecifying the distributions of the priors, Baltagi et al. (2018) considered the ε-contamination class of prior distributions for \((\beta, b, \tau)\):

\[
\Gamma = \{ \pi(\beta, b, \tau|g_0, h_0) = (1 - \varepsilon) \pi_0(\beta, b, \tau|g_0, h_0) + \varepsilon q(\beta, b, \tau|g_0, h_0) \},
\]

where \(\pi_0(\cdot)\) is the base elicited prior, \(q(\cdot)\) is the contamination belonging to some suitable class \(Q\) of prior distributions, and \(0 \leq \varepsilon \leq 1\) reflects the amount of error in \(\pi_0(\cdot)\). \(\tau\) is assumed to have a vague prior, \(p(\tau) \propto \tau^{-1}\), \(0 < \tau < \infty\), and \(\pi_0(\beta, b, \tau|g_0, h_0)\) is the base prior assumed to be a specific \(g\)-prior with

\[
\begin{align*}
\beta & \sim N(b_{0tK_x}, (\tau g_0 \Lambda_X)^{-1}) \quad \text{with} \quad \Lambda_X = X'X, \\
\quad b & \sim N(b_{0tK_x}, (\tau h_0 \Lambda_W)^{-1}) \quad \text{with} \quad \Lambda_W = W'W, 
\end{align*}
\]

where \(t_{K_x}\) is a \((K_x \times 1)\) vector of ones. Here, \(\beta_0, b_0, g_0\) and \(h_0\) are known scalar hyperparameters of the base prior \(\pi_0(\beta, b, \tau|g_0, h_0)\). The probability density function (henceforth pdf) of the base prior \(\pi_0(\cdot)\) is given by:

\[
\pi_0(\beta, b, \tau|g_0, h_0) = p(\beta|b, \tau, \beta_0, b_0, g_0, h_0) \times p(b|\tau, b_0, h_0) \times p(\tau). 
\]

The possible class of contamination \(Q\) is defined as:

\[
Q = \left\{ q(\beta, b, \tau|g_0, h_0) = p(\beta|b, \tau, \beta_q, b_q, g_q, h_q) \times p(b|\tau, b_q, h_q) \times p(\tau) \quad \text{with} \quad 0 < g_q \leq g_0, \quad 0 < h_q \leq h_0 \right\},
\]

with

\[
\begin{align*}
\beta & \sim N(b_{qK_x}, (\tau g_q \Lambda_X)^{-1}), \\
\quad b & \sim N(b_{qK_x}, (\tau h_q \Lambda_W)^{-1}), 
\end{align*}
\]

where \(\beta_q, b_q, g_q\) and \(h_q\) are unknown. The restrictions \(g_q \leq g_0\) and \(h_q \leq h_0\) imply that the base prior is the best possible so that the precision of the base prior is greater than any prior belonging to the contamination class. The ε-contamination class of prior distributions for \((\beta, b, \tau)\) is then conditional on known \(g_0\) and \(h_0\).

Following Baltagi et al. (2018) for the static panel model, we use a two-step strategy because it simplifies the derivation of the predictive densities (or marginal likelihoods).\(^3\) This will be extended to the dynamic panel model introduced in the next section.

### 2.2. The dynamic framework

This paper considers the Gaussian dynamic linear mixed model:

\[
y_{it} = \rho y_{it-1} + X'_{it}\beta + W'_{it}b_i + u_{it} = Z'_{it}\theta + W'_{it}b_i + u_{it}, \quad i = 1, ..., N, \quad t = 2, ..., T, 
\]

\(^3\) One could also use a one-step estimation of the ML-II posterior distribution. But in the one-step approach, the pdf of \(y\) and the pdf of the base prior \(\pi_0(\beta, b, \tau|g_0, h_0)\) need to be combined to get the predictive density. It thus leads to a complex expression whose integration with respect to \((\beta, b, \tau)\) may be involved.
where $Z_t' = [y_{it-1}, X_t']'$ and $\theta' = [\rho, \beta']'$ is a $(1 \times K_1)$ vector with $K_1 = K_y + 1$. The likelihood is conditional on the first observations $y_1$. In that case, the first period is assumed exogenous and known. In the spirit of eq(2), we have the following:

First stage: 
$$y = \rho y_{t-1} + X\beta + Wb + u, \ u \sim N(0, \Sigma), \Sigma = \tau^{-1} I_{N(T-1)}$$

Second stage: 
$$\beta \sim N(\beta_0, \Lambda_\beta) \ \text{and} \ b \sim N(b_0, \Lambda_b)$$

with 
$$p(\tau) \propto \tau^{-1}, \ \Lambda_\beta = (\tau g'X'X)^{-1} \ \text{and} \ \Lambda_b = (\tau hW'W)^{-1}.$$ 

where $y = (y_{1,1}, ..., y_{1,T}, ..., y_{N,2}, ..., y_{N,T})'$ and $y_{t-1} = (y_{1,1}, ..., y_{1,T-1}, ..., y_{N,1}, ..., y_{N,T-1})'$ are $(N(T - 1) \times 1)$. $X$ is $(N(T - 1) \times K_y)$, $W$ is $(N(T - 1) \times K_z)$, $u$ is $(N(T - 1) \times 1)$ and $I_{N(T-1)}$ is a $(N(T - 1) \times N(T - 1))$ identity matrix.

There is an extensive literature on autoregressive processes using Bayesian methods. The stationarity assumption implies that the autoregressive time dependence parameter space for $\rho$ is a compact subset of $(-1, 1)$. For the pros and cons of imposing a stationarity hypothesis in a Bayesian setup see Phillips (1991). Ghosh and Heo (2003) proposed a comparative study using some selected noninformative (objective) priors for the AR(1) model. Ibazizen and Fellag (2003), assumed a noninformative prior for the autoregressive parameter without considering the stationarity assumption for the AR(1) model. However, most papers consider a noninformative (objective) prior for the Bayesian analysis of an AR(1) model without considering the stationarity assumption. See for example DeJong and Whiteman (1991), Schotman and van Dijk (1991), and Sims and Uhlig (1991). For the dynamic random coefficients panel data model, Hsiao and Pesaran (2008) do not impose any constraint on the coefficients of the lag dependent variable, $\rho_t$. But, following Liu and Tiao (1980), they suggest that one way to impose the stability condition on individual units would be to assume that $\rho_t$ follows a rescaled Beta distribution on $(0, 1)$. In the time series framework, and for an AR(1) model, Karakani et al. (2016) have performed a posterior sensitivity analysis based on Gibbs sampling with four different priors: natural conjugate prior, Jeffreys’ prior, truncated normal prior and $g$-prior. Their respective performances are compared in terms of the highest posterior density region criterion. They show that the truncated normal distribution outperforms very slightly the $g$-prior and more strongly the other priors especially when the time dimension is small. On the other hand, for a larger time span, there is no significant difference between the truncated normal distribution and the $g$-prior.

Nevertheless, introducing a truncated normal distribution for $\rho$ poses very complex integration problems due to the presence of the normal cdf function as integrand in the marginal likelihoods with $\varepsilon$-contamination class of prior distributions. To avoid these problems, $\rho$ is assumed to be $U(-1, 1)$. In that case, its mean (0) and its variance (1/3) are exactly defined and we do not need to introduce an $\varepsilon$-contamination class of prior distributions for $\rho$ at the second stage of the hierarchy. This was initially our first goal (see appendix A.1 in the supplementary material). Unfortunately, the results using Monte Carlo simulations showed biased estimates of $\rho$, $\beta$ and residual variances (see Appendix A.2 in the supplementary material). Consequently, we assume a Zellner $g$-prior, for $\theta' = [\rho, \beta']'$ which encompasses the coefficient of the lagged dependent variable $y_{it-1}$ and those of the explanatory variables $X_{it}'$. In other words, the two-stage hierarchy becomes:

First stage: 
$$y = Z_{it}'\theta + Wb + u, \ u \sim N(0, \Sigma), \Sigma = \tau^{-1} I_{N(T-1)}$$

Second stage: 
$$\theta \sim N(\theta_0, \Lambda_\theta) \ \text{and} \ b \sim N(b_0, \Lambda_b)$$

with 
$$p(\tau) \propto \tau^{-1}, \ \Lambda_\theta = (\tau gZ'Z)^{-1} \ \text{and} \ \Lambda_b = (\tau hW'W)^{-1}.$$
Thus, we do not impose stationarity constraints like many authors and we respect the philosophy of \( \varepsilon \)-contamination class using data-driven priors.

2.3. The robust dynamic linear model in the two-stage hierarchy

Using a two-step approach, we can integrate first with respect to \((\theta, \tau)\) given \(b\) and then, conditional on \(\theta\), we integrate with respect to \((b, \tau)\).

1. Let \(y^* = (y - Wb)\). Derive the conditional ML-II posterior distribution of \(\theta\) given the specific effects \(b\).

2. Let \(\tilde{y} = (y - Z\theta)\). Derive the conditional ML-II posterior distribution of \(b\) given the coefficients \(\theta\).

Thus, the marginal likelihoods (or predictive densities) corresponding to the base priors are:

\[
m(y^*|\pi_0, b, g_0) = \int_0^\infty \int_{\mathbb{R}^{k_1}} \pi_0(\theta, \tau|g_0) \times p(y^*|Z, b, \tau) \ d\theta \ d\tau
\]

and

\[
m(\tilde{y}|\pi_0, \theta, h_0) = \int_0^\infty \int_{\mathbb{R}^{k_2}} \pi_0(b, \tau|h_0) \times p(\tilde{y}|W, \theta, \tau) \ db \ d\tau,
\]

with

\[
\pi_0(\theta, \tau|g_0) = \frac{(\tau g_0)^{\frac{k_1}{2}}}{(2\pi)^{\frac{k_1}{2}}} \tau^{-1} |\Lambda_Z|^{1/2} \exp \left( -\frac{\tau g_0}{2} (\theta - \theta_0_1 K_1)' \Lambda_Z (\theta - \theta_0_1 K_1) \right),
\]

\[
\pi_0(b, \tau|h_0) = \frac{(\tau h_0)^{\frac{k_2}{2}}}{(2\pi)^{\frac{k_2}{2}}} \tau^{-1} |\Lambda_W|^{1/2} \exp \left( -\frac{\tau h_0}{2} (b - b_0_1 K_2)' \Lambda_W (b - b_0_1 K_2) \right).
\]

Solving these equations is considerably easier than solving the equivalent expression in the one-step approach.

2.3.1. The first step of the robust Bayesian estimator

Let \(y^* = y - Wb\). Combining the pdf of \(y^*\) and the pdf of the base prior, we get the predictive density corresponding to the base prior\(^4\):

\[
m(y^*|\pi_0, b, g_0) = \int_0^\infty \int_{\mathbb{R}^{k_1}} \pi_0(\theta, \tau|g_0) \times p(y^*|Z, b, \tau) \ d\theta \ d\tau
\]

\[
= \tilde{H} \left( \frac{g_0}{g_0 + 1} \right)^{K_1/2} \left( 1 + \left( \frac{g_0}{g_0 + 1} \right) \left( \frac{R_0^2}{1 - R_0^2} \right) \right)^{-\frac{NT}{2}}
\]

with \(\tilde{H} = \frac{\Gamma\left(\frac{NT}{2}\right)}{\pi\left(\frac{2\pi}{v(b)}\right)^{\frac{NT}{2}}}\), \(R_0^2 = \frac{(\tilde{\theta}(b) - \theta_0_1 K_1)' \Lambda_Z (\tilde{\theta}(b) - \theta_0_1 K_1)}{(\theta(b) - \theta_0_1 K_1)' \Lambda_Z (\theta(b) - \theta_0_1 K_1) + v(b)}\), \(\tilde{\theta}(b) = \Lambda_Z^{-1} Z' y^*\) and \(v(b) = (y^* - Z\tilde{\theta}(b))'(y^* - Z\tilde{\theta}(b))\), and where \(\Gamma(\cdot)\) is the Gamma function.

---

\(^4\)Derivation can be found in the supplementary appendix of Baltagi et al. (2018).
Likewise, we can obtain the predictive density corresponding to the contaminated prior for the distribution \( q(\theta, \tau | g_0, h_0) \in Q \) from the class \( Q \) of possible contamination distributions:

\[
m(y^* | q, b, g_0) = \tilde{H} \left( \frac{g_0}{g_0 + 1} \right)^{\frac{\kappa_y}{2}} \left( 1 + \left( \frac{g_0}{g_0 + 1} \right) \left( \frac{R_{\theta_q}^2}{1 - R_{\theta_q}^2} \right) \right)^{-\frac{\lambda_y}{2}}, \tag{12}
\]

where

\[
R_{\theta_q}^2 = \frac{(\hat{\theta}(b) - \theta_q \ell K_1)\Lambda Z(\hat{\theta}(b) - \theta_q \ell K_1)}{(\hat{\theta}(b) - \theta_q \ell K_1)\Lambda Z(\hat{\theta}(b) - \theta_q \ell K_1) + v(b)}.
\]

As the \( \varepsilon \)-contamination of the prior distributions for \((\theta, \tau)\) is defined by \( \pi(\theta, \tau | g_0) = (1 - \varepsilon) \pi_0(\theta, \tau | g_0) + \varepsilon q(\theta, \tau | g_0) \), the corresponding predictive density is given by:

\[
m(y^* | \pi, b, g_0) = (1 - \varepsilon) m(y^* | \pi_0, b, g_0) + \varepsilon m(y^* | q, b, g_0)
\]

and

\[
\sup_{\pi \in \Gamma} m(y^* | \pi, b, g_0) = (1 - \varepsilon) m(y^* | \pi_0, b, g_0) + \varepsilon \sup_{q \in Q} m(y^* | q, b, g_0).
\]

The maximization of \( m(y^* | \pi, b, g_0) \) requires the maximization of \( m(y^* | q, b, g_0) \) with respect to \( \theta_q \) and \( g_0 \). The first-order conditions lead to

\[
\hat{\theta}_q = (\ell K_1 \Lambda Z \ell K_1)^{-1} \ell K_1 \Lambda Z \hat{\theta}(b) \tag{13}
\]

and

\[
\hat{g}_q = \min(g_0, g^*),
\]

with \( g^* = \max \left[ \left( \frac{(NT - K_1)(\hat{\theta}(b) - \theta_q \ell K_1)\Lambda Z(\hat{\theta}(b) - \theta_q \ell K_1)}{v(b)} - 1 \right)^{-1}, 0 \right] \]

\[
= \max \left[ \left( \frac{(NT - K_1)(R_{\theta_q}^2)}{1 - R_{\theta_q}^2} - 1 \right)^{-1}, 0 \right].
\]

Denote \( \sup_{q \in Q} m(y^* | q, b, g_0) = m(y^* | \tilde{q}, b, g_0) \). Then

\[
m(y^* | \tilde{q}, b, g_0) = \tilde{H} \left( \frac{\tilde{g}_q}{\tilde{g}_q + 1} \right)^{\frac{\kappa_y}{2}} \left( 1 + \left( \frac{\tilde{g}_q}{\tilde{g}_q + 1} \right) \left( \frac{R_{\theta_q}^2}{1 - R_{\theta_q}^2} \right) \right)^{-\frac{\lambda_y}{2}}.
\]

Let \( \pi_0^*(\theta, \tau | g_0) \) denote the posterior density of \((\theta, \tau)\) based upon the prior \( \pi_0(\theta, \tau | g_0) \). Also, let \( q^*(\theta, \tau | g_0) \) denote the posterior density of \((\theta, \tau)\) based upon the prior \( q(\theta, \tau | g_0) \). The ML-II posterior density of \( \theta \) is thus given by:

\[
\hat{\pi}^*(\theta | g_0) = \int_0^\infty \hat{\pi}^*(\theta, \tau | g_0) \, d\tau
\]

\[
= \hat{\lambda}_{\theta,g_0} \int_0^\infty \pi_0^*(\theta, \tau | g_0) \, d\tau + \left( 1 - \hat{\lambda}_{\theta,g_0} \right) \int_0^\infty q^*(\theta, \tau | g_0) \, d\tau
\]

\[
= \hat{\lambda}_{\theta,g_0} \pi_0^*(\theta | g_0) + \left( 1 - \hat{\lambda}_{\theta,g_0} \right) \hat{q}^*(\theta | g_0) \tag{15}
\]
The ML-II posterior density of \( \theta \). Note that \( \lambda_{\theta, g_0} \) depends upon the ratio of the \( R_{\theta, g_0}^2 \) and \( R_{\theta_0}^2 \) but primarily on the sample size \( NT \). Indeed, \( \lambda_{\theta, g_0} \) tends to 0 when \( R_{\theta, g_0}^2 > R_{\theta_0}^2 \) and tends to 1 when \( R_{\theta, g_0}^2 < R_{\theta_0}^2 \), irrespective of the model fit (i.e., the absolute values of \( R_{\theta, g_0}^2 \) or \( R_{\theta_0}^2 \)). Only the relative values of \( R_{\theta, g_0}^2 \) and \( R_{\theta_0}^2 \) matter.

It can be shown that \( \pi_0^* (\theta \mid g_0) \) is the pdf (see the supplementary appendix of Baltagi et al. (2018)) of a multivariate \( t \)-distribution with mean vector \( \theta_\ast (b \mid g_0) \), variance-covariance matrix \( \left( \frac{\xi_{q, \theta} M_{q, \theta}}{NT-2} \right) \) and degrees of freedom (NT) with

\[
M_{0, \theta} = \frac{(g_0 + 1)}{v(b)} \Lambda_Z \text{ and } \xi_{0, \theta} = 1 + \frac{-g_0}{(g_0 + 1)} \left( \frac{R_{\theta, g_0}^2}{1 - R_{\theta_0}^2} \right).
\]

(16)

\( \theta_\ast (b \mid g_0) \) is the Bayes estimate of \( \theta \) for the prior distribution \( \pi_0 (\theta, \tau) \):

\[
\theta_\ast (b \mid g_0) = \frac{\hat{\theta} (b) + g_0 \theta_0 \epsilon_{K, 1}}{g_0 + 1}.
\]

(17)

Likewise \( \pi_\ast (\theta \mid g_0) \) is the pdf of a multivariate \( t \)-distribution with mean vector \( \hat{\theta}_{EB} (b \mid g_0) \), variance-covariance matrix \( \left( \frac{\xi_{q, \theta} M_{q, \theta}}{NT-2} \right) \) and degrees of freedom (NT) with

\[
\xi_{q, \theta} = 1 + \frac{-\hat{g}_q}{(\hat{g}_q + 1)} \left( \frac{R_{\hat{\theta}, q_0}^2}{1 - R_{\theta_0}^2} \right) \text{ and } M_{q, \theta} = \left( \frac{(\hat{g}_q + 1)}{v(b)} \right) \Lambda_Z,
\]

(18)

where \( \hat{\theta}_{EB} (b \mid g_0) \) is the empirical Bayes estimator of \( \theta \) for the contaminated prior distribution \( q (\theta, \tau) \) given by:

\[
\hat{\theta}_{EB} (b \mid g_0) = \frac{\hat{\theta} (b) + \hat{\theta}_q \epsilon_{K_1}}{\hat{g}_q + 1}.
\]

(19)

The mean of the ML-II posterior density of \( \theta \) is then:

\[
\hat{\theta}_{ML-II} = E [\pi_\ast (\theta \mid g_0)] = \hat{\lambda}_{\theta, g_0} E [\pi_0^* (\theta \mid g_0)] + \left( 1 - \hat{\lambda}_{\theta, g_0} \right) E [\pi_\ast (\theta \mid g_0)]
\]

\[
= \hat{\lambda}_{\theta, g_0} \theta_\ast (b \mid g_0) + \left( 1 - \hat{\lambda}_{\theta, g_0} \right) \hat{\theta}_{EB} (b \mid g_0).
\]

The ML-II posterior density of \( \theta \), given \( b \) and \( g_0 \) is a shrinkage estimator. It is a weighted average of the Bayes estimator \( \theta_\ast (b \mid g_0) \) under base prior \( g_0 \) and the data-dependent empirical Bayes estimator \( \hat{\theta}_{EB} (b \mid g_0) \). If the base prior is consistent with the data, the weight \( \hat{\lambda}_{\theta, g_0} \to 1 \) and the ML-II posterior density of \( \theta \) gives more weight to the posterior \( \pi_0^* (\theta \mid g_0) \) derived from the elicited prior.
where $\theta$ and matrix $\lambda$.

2.3.2. The second step of the robust Bayesian estimator

In this case $\hat{\theta}_{ML-II}$ is close to the Bayes estimator $\theta_*(b|g_0)$. Conversely, if the base prior is not consistent with the data, the weight $\hat{\lambda}_{g_0} \to 0$ and the ML-II posterior density of $\theta$ is then close to the posterior $\tilde{q}^*(\theta|g_0)$ and to the empirical Bayes estimator $\hat{\theta}_{EB} (b|g_0)$. The ability of the $\varepsilon$-contamination model to extract more information from the data is what makes it superior to the classical Bayes estimator based on a single base prior.\(^5\)

Let $\tilde{y} = y - Z\theta$. Moving along the lines of the first step, the ML-II posterior density of $b$ is given by:

$$\pi^* (b|h_0) = \tilde{\lambda}_{b,h_0}\pi_0^* (b|h_0) + \left(1 - \tilde{\lambda}_{b,h_0}\right) \tilde{q}^* (b|h_0)$$

with

$$\tilde{\lambda}_{b,h_0} = \left[1 + \frac{\varepsilon}{1 - \varepsilon} \left(\frac{\tilde{h}}{h_0 + 1} \frac{h_0}{h_0 + 1}\right)^{K_2/2} \left(1 + \left(h_0 T_{\varepsilon}\right) \left(\frac{R_0}{1 - R_0}\right)^{\frac{1}{2}}\right)^{-1}\right],$$

where

$$R_{\theta_0}^2 = \frac{(b(\theta) - b_0 t_{K_2})'\Lambda W (b(\theta) - b_0 t_{K_2})}{(b(\theta) - b_0 t_{K_2})'\Lambda W (b(\theta) - b_0 t_{K_2}) + v(\theta)},$$

$$R_{\theta_0}^2 = \frac{(\tilde{b}(\theta) - \tilde{b}_0 t_{K_2})'\Lambda W (\tilde{b}(\theta) - \tilde{b}_0 t_{K_2})}{(\tilde{b}(\theta) - \tilde{b}_0 t_{K_2})'\Lambda W (\tilde{b}(\theta) - \tilde{b}_0 t_{K_2}) + v(\theta)},$$

with $\tilde{b}(\theta) = \Lambda_W^{-1} W' \tilde{y}$ and $v(\theta) = (\tilde{y} - W\tilde{b}(\theta))'(\tilde{y} - W\tilde{b})(\theta)$,

$$\tilde{b}_q = \left(\Lambda_W^{-1} W' K_2\right)^{-1} \Lambda_W \tilde{b}(\theta)$$

and

$$\hat{\lambda}_q = \min (h_0, h^*)$$

with $h^* = \max \left[\left(\frac{(NT - K_2) (\tilde{b}(\theta) - \tilde{b}_q t_{K_2})'\Lambda_W (\tilde{b}(\theta) - \tilde{b}_q t_{K_2})}{v(\theta)} - 1\right)^{-1}, 0\right]$

$$= \max \left[\left(\frac{(NT - K_2)}{K_2} \left(\frac{R_{\theta_0}^2}{1 - R_{\theta_0}^2}\right) - 1\right)^{-1}, 0\right].$$

$\pi_0^* (b|h_0)$ is the pdf of a multivariate $t$-distribution with mean vector $b_*(\theta|h_0)$, variance-covariance matrix $\left(\frac{\xi_0, h_0, M_0, b^2}{NT - 2}\right)$ and degrees of freedom $(NT)$ with

$$M_0, b = \frac{(h_0 + 1)}{v(\theta)} \Lambda_W$$

and $\xi_0, b = 1 + \left(\frac{h_0}{h_0 + 1}\right) \frac{(\tilde{b}(\theta) - b_0 t_{K_2})'\Lambda_W (\tilde{b}(\theta) - b_0 t_{K_2})}{v(\theta)}$.

\(^5\)Following Berger (1985), Baltagi et al. (2018) derived the analytical ML-II posterior variance-covariance matrix of $\theta$ (see the supplementary appendix of Baltagi et al. (2018)).
\( b_*(\theta|h_0) \) is the Bayes estimate of \( b \) for the prior distribution \( \pi_0 (b, \tau|h_0) : \\
\hat{b}_*(\theta|h_0) = \frac{\hat{b}(\theta) + h_0 b_0 \kappa_2}{h_0 + 1}. \\
q^* (b|h_0) \) is the pdf of a multivariate t-distribution with mean vector \( \hat{b}_{EB} (\theta|h_0) \), variance-covariance matrix \( \left( \frac{\xi_{1,b} M_{1,b}}{N_{b}} \right) \) and degrees of freedom \( (NT) \) with \( \xi_{1,b} = 1 + \left( \frac{\hat{h}_q}{h_q + 1} \right) \left( \frac{\hat{b}(\theta) - \hat{b}_{EB} (\theta) - \hat{b}_{q} \kappa_2}{v(\theta)} \right) \Lambda_W \) and \( M_{1,b} = \left( \frac{\hat{h}_q + 1}{v(\theta)} \right) \Lambda_W \). \\
\hat{b}_{EB} (\theta|h_0) \) is the empirical Bayes estimator of \( b \) for the contaminated prior distribution \( q (b, \tau|h_0) : \\
\hat{b}_{EB} (\theta|h_0) = \hat{\theta}(b) + \hat{h}_q \hat{b}_{q} \kappa_2. \\
The mean of the ML-II posterior density of \( b \) is hence given by: \\
\bar{b}_{ML-II} = \hat{\lambda}_b b_*(\theta|h_0) + \left( 1 - \hat{\lambda}_b \right) \hat{b}_{EB} (\theta|h_0). (21) \\
The ML-II posterior variance-covariance matrix of \( b \) can be derived in a similar fashion to that of \( \hat{\theta}_{ML-II} \).

2.4. Estimating the ML-II posterior variance-covariance matrix

Many have raised concerns about the unbiasedness of the posterior variance-covariance matrices of \( \hat{\theta}_{ML-II} \) and \( b_{ML-II} \). Indeed, they will both be biased towards zero as \( \hat{\lambda}_{\theta,g_0} \) and \( \hat{\lambda}_{b,h_0} \) go to zero and converge to the empirical variance which is known to underestimate the true variance (see e.g. Berger and Berliner (1986); Gilks et al. (1997); Robert (2007)). Consequently, the assessment of the performance of either \( \hat{\theta}_{ML-II} \) or \( \hat{b}_{ML-II} \) using standard quadratic loss functions can not be conducted using the analytical expressions. What is needed is an unbiased estimator of the true ML-II variances. Baltagi et al. (2018) proposed two different strategies to approximate these, each with different desirable properties: MCMC with multivariate t-distributions or block resampling bootstrap. Simulations show that one needs as few as 20 bootstrap samples to achieve acceptable results. Here, we will use the same individual block resampling bootstrap method. Following Bellman et al. (1989); Andersson and Karlsson (2001), and Kapetanios (2008), individual block resampling consists of drawing an \( (N \times T) \) matrix \( Y^{BR} \) whose rows are obtained by resampling those of an \( (N \times T) \) matrix \( Y \) with replacement. Conditionally on \( Y \), the rows of \( Y^{BR} \) are independent and identically distributed. The following algorithm is used to approximate the variance matrices:

1. Loop over \( BR \) samples

---

6See the supplementary appendix of Baltagi et al. (2018).

7Increasing the number of bootstrap samples does not change the results but increases the computation time considerably.
2. In the first step, compute the mean of the ML-II posterior density of \( \theta \) using our initial shrinkage procedure

\[
\hat{\theta}_{ML-II,br} = E[\hat{\pi}^*(\theta|g_0)] = \lambda_{\theta,g_0} \theta_0 + \left(1 - \lambda_{\theta,g_0}\right) \hat{\theta}_{EB}(b|g_0).
\]

3. In the second step, compute the mean of the ML-II posterior density of \( b \):

\[
\hat{b}_{ML-II,br} = \lambda_{b,b} \theta_0 + \left(1 - \lambda_{b,b}\right) \hat{b}_{EB}(\theta|g_0).
\]

4. Once the \( BR \) bootstraps are completed, use the \( (K_1 \times BR) \) matrix of coefficients \( \theta^{(BR)} \) and the \( (N \times BR) \) matrix of coefficients \( b^{(BR)} \) to compute:

\[
\hat{\theta}_{ML-II} = E[\theta^{(BR)}], \quad \hat{\sigma}_{\theta,ML-II} = \sqrt{\text{diag}(\text{Var}[\theta^{(BR)}])},
\]

\[
\hat{b}_{ML-II} = E[b^{(BR)}], \quad \hat{\sigma}_{b,ML-II} = \sqrt{\text{diag}(\text{Var}[b^{(BR)}])}.
\]

3. Monte Carlo simulation study

In what follows, we compare the finite sample properties of our proposed estimator with those of standard classical estimators.

3.1. The DGP of the Monte Carlo simulation study

For the random effects (RE) world, we use the same DGP as that of Baltagi et al. (2018) extended to the dynamic case. For the dynamic homogeneous/heterogeneous panel data model with common trends or with common correlated effects, we will follow Chudik and Pesaran (2015a,b).

\[
y_{it} = \rho y_{i,t-1} + x_{1,1,1,1} + x_{1,2,1,1} + x_{1,2,1,2} + x_{2,1,1,2} + x_{2,2,1,2} + \eta_1 + \eta_2 + \mu_i + u_{it}, \quad (22)
\]

for \( i = 1, \ldots, N \), \( t = 2, \ldots, T \), with

\[
x_{1,1,1,1} = 0.7x_{1,1,1,t-1} + \delta_{i} + \zeta_{it}
\]

\[
x_{1,2,1,1} = 0.7x_{1,2,1,t-1} + \theta_{i} + \zeta_{it}
\]

\[
u_{it} \sim N(0, \tau^{-1}), \quad \left(\delta_{i} + \theta_{i} + \zeta_{it}, \varsigma_{it}\right) \sim U(-2,2)
\]

and \( \rho = 0.75, \beta_{1,1} = \beta_{1,2} = \beta_{2} = 1. \)

1. For a random effects (RE) world, we assume that:

\[
\eta_1 = \eta_2 = 0
\]

\[
x_{2,1,1,2} = 0.7x_{2,1,1,t-1} + \kappa_{i} + \vartheta_{i,1} + \psi_{i,1} \sim U(-2,2)
\]

\[
\mu_i \sim N(0, \sigma_{\mu}^2) \quad \sigma_{\mu}^2 = 4\tau^{-1}.
\]

Furthermore, \( x_{1,1,1,1}, x_{1,2,1,2} \) and \( x_{2,1,1,2} \) are assumed to be exogenous in that they are not correlated with \( \mu_i \) and \( u_{it} \).
2. For a Chamberlain-type fixed effects (FE) world, we assume that:

\[ \eta_1 = \eta_2 = 0; \]
\[ x_{2,it} = \delta_{2,i} + \omega_{2,it} , \delta_{2,i} \sim N(m_{\delta_2}, \sigma_{\delta_2}^2), \omega_{2,it} \sim N(m_{\omega_2}, \sigma_{\omega_2}^2); \]
\[ m_{\delta_2} = m_{\omega_2} = 1, \sigma_{\delta_2}^2 = 8, \sigma_{\omega_2}^2 = 2; \]
\[ \mu_i = x_{2,1i} + x_{2,2i} \pi + \ldots + x_{2,IT} \pi_T + \nu_i, \nu_i \sim N(0, \sigma_{\nu_i}^2); \]
\[ \sigma_{\nu_i}^2 = 1, \pi_t = (0.8)^{T-t} \text{ for } t = 1, \ldots, T. \]

\( x_{1,1,it} \) and \( x_{1,2,it} \) are assumed to be exogenous but \( x_{2,it} \) is correlated with the \( \mu_i \) and we assume an exponential growth for the correlation coefficient \( \pi_t \).

3. For a Hausman-Taylor (HT) world, we assume that:

\[ \eta_1 = \eta_2 = 1; \]
\[ x_{2,it} = 0.7x_{2,it-1} + \mu_i + \vartheta_{it} , \vartheta_{it} \sim U(-2, 2); \]
\[ V_{1,i} = 1, \forall i; \]
\[ V_{2,i} = \mu_i + \delta_i + \theta_i + \xi_i, \xi_i \sim U(-2, 2); \]
\[ \mu_i \sim N(0, \sigma_{\mu_i}^2) \text{ and } \sigma_{\mu_i}^2 = 4 \pi^{-1}. \]

\( x_{1,1,it} \) and \( x_{1,2,it} \) and \( V_{1,i} \) are assumed to be exogenous while \( x_{2,it} \) and \( V_{2,i} \) are endogenous because they are correlated with the \( \mu_i \) but not with the \( u_{it} \).

4. For the homogeneous panel data world with common trends, we follow Chudik and Pesaran (2015a,b) and assume that

\[ y_{it} = \rho y_{it-1} + x_{it} \beta_1 + x_{it-1} \beta_2 + f^\prime_i \gamma_i + u_{it}, \text{ for } i = 1, \ldots, N, \ t = 2, \ldots, T, \quad (23) \]

with

\[ x_{it} = \alpha_x y_{it-1} + f^\prime_i \gamma_i + \omega_{x_{it}} \]
\[ \omega_{x_{it}} = \varphi_x \omega_{x-it-1} + \zeta_{x_{it}} \]
\[ \gamma_{l} = \gamma_l + \eta_l \gamma_i, \text{ for } l = 1, \ldots, m \]
\[ \gamma_{x_{it}} = \gamma_{x_{l}} + \eta_l \gamma_{x_{l}}, \text{ for } l = 1, \ldots, m \]

where

\[ \zeta_{x_{it}} \sim N(0, \sigma_{\zeta_{x_{it}}}^2) , \quad \sigma_{\zeta_{x_{it}}}^2 = \beta_1 \left[ 1 - |E(\varphi_x)| \right] = \varphi_{x_{it}} \sim N(0, 0.9), \text{ for } i = 1, \ldots, N \]
\[ \eta_l \gamma_i \sim N(0, \sigma_{\eta_l \gamma_i}^2) , \quad \eta_l \gamma_{x_{l}} \sim N(0, \sigma_{\eta_l \gamma_{x_{l}}}^2), \text{ for } l = 1, \ldots, m \]
\[ \gamma_l = \sqrt{T} \times c_{\gamma_l}, \quad \gamma_{x_{l}} = \sqrt{T} \times c_{x_{l}} \]
\[ c_{x_{l}} = \frac{2}{m(m+1)} - \frac{2 \sigma_{x_{l}}^2}{(m+1)}, \quad \rho = 0.75, \beta_1 = \beta_2 = 1 \]
\[ \text{and } u_{it} \sim N(0, \tau^{-1}). \]

\( f_i \) and \( \gamma_i \) are \((m \times 1)\) vectors. We consider \( m = 2 \) deterministic known common trends: one linear trend \( f_{t,1} = t/T \) and one polynomial trend: \( f_{t,2} = t/T + 1.4(t/T)^2 - 3(t/T)^3 \) for \( t = 1, \ldots, T \). The feedback coefficients follow a uniform distribution \( \alpha_{x_{l}} \sim U(0, 0.15) \) and are non-zero for all \( i (\alpha_{x_{l}} \neq 0) \). They lead to weakly exogenous regressors \( x_{it} \).
5. For the homogeneous panel data world with correlated common effects, we follow Chudik and Pesaran (2015a,b), and assume that the m common trends, \( f_l \) (23), are replaced with unobserved common factors:

\[
fu = \rho f_{l-1} x + \xi fu, \quad \xi fu \sim N(0, 1 - \rho^2_{fu}), \quad l = 1, ..., m.
\]

We assume that the common factors are independent stationary AR(1) processes with \( \rho_{fu} = 0.6 \) for \( l = 1, ..., m \).

6. For the heterogeneous panel data world with correlated common effects, we follow Chudik and Pesaran (2015a,b) and assume that \( \rho \) (resp. \( \beta_l \)) in the model (23) is replaced by individual coefficients \( \rho_i \sim U(0.6, 0.9) \) (resp. \( \beta_{li} \sim U(0.5, 1) \)) for \( i = 1, ..., N \) and we keep the \( m \) unobserved common factors as defined previously.

For each set-up, we vary the size of the sample and the length of the panel. We choose several \((N, T)\) pairs with \( N = 100, 200 \) and \( T = 10, 30 \) for cases 1 to 3 and \( N = (50, 100) \) and \( T = (30, 50) \) for cases 4 to 6. The autoregressive coefficient is set as \( \rho = 0.75 \). We set the initial values of \( y_{it}, x_{1,1,it}, x_{1,2,it} \) and \( x_{2,it} \) to zero. We next generate all the \( x_{1,1,it}, x_{1,2,it}, x_{2,it}, y_{it}, u_{it}, \xi_{it}, \Delta_{it}, \omega_{2,it}, ... \) over \( T + T_0 \) time periods and we drop the first \( T_0 \) observations to reduce the dependence on the initial values. The robust Bayesian estimators for the two-stage hierarchy are estimated with \( \varepsilon = 0.5 \), though we investigate the robustness of our results to various values of \( \varepsilon \).

We must set the hyperparameters values \( \theta_0, b_0, g_0, a_0, \tau \) for the initial distributions of \( \theta \sim N(\theta_{0,K_1}, (\tau g_0 A_2)^{-1}) \) and \( b \sim N(b_{0,K_2}, (\tau h_0 A_W)^{-1}) \) where \( \theta = [\rho, \beta_{1,1}, \beta_{1,2}, \beta_2]^T \) for the first three cases and \( \theta = [\rho, \beta_1, \beta_2]^T \) for the last three cases. While we can choose arbitrary values for \( \theta_0, b_0 \) and \( \tau \), the literature generally recommends using the regular unit information prior (UIP) to set the \( g \)-priors. In the normal regression case, and following Kass and Wasserman (1995), the UIP corresponds to \( g_0 = h_0 = 1/NT \), leading to Bayes factors that behave like the Bayesian Information Criterion (BIC).

For the 28 robust estimators, we use \( BR = 20 \) samples in the block resampling bootstrap. For each experiment, we run \( R = 1,000 \) replications and we compute the means, standard errors and root mean squared errors (RMSEs) of the coefficients and the residual variances.

3.1.1. The random effects world

Rewrite the general dynamic model (8) as follows:

\[
y = Z0 + Wb + u = Z0 + Z\mu + u
\]

with \( Z_{it} = [y_{it}, x_{1,1,it}, x_{1,2,it}] \), \( \theta = [\rho, \beta] \) and \( X_{it} = [x_{1,1,it}, x_{1,2,it}, x_{2,it}] \),

where \( u \sim N(0, \Sigma), \Sigma = \tau^{-1}I_{N(T-1)}, Z\mu = I_N \otimes \iota_{T-1} \) is \((N(T - 1) \times N)\), \( \otimes \) is the Kronecker product, \( \iota_{T-1} \) is a \((T - 1 \times 1)\) vector of ones and \( \mu(\equiv b) \) is a \((N \times 1)\) vector of idiosyncratic parameters. When \( W \equiv Z\mu \), the random effects, \( \mu \sim N(0, \sigma^2\mu I_N) \), are associated with the error term \( \nu = Z\mu + u \) with \( \text{Var}(\nu) = \sigma^2\mu (I_N \otimes J_{T-1}) + \sigma^2\nu I_{N(T-1)} \), where \( J_{T-1} = \iota_{T-1}^T\iota_{T-1} \).

---

8\( \varepsilon = 0.5 \) is an arbitrary value. This implicitly assumes that the amount of error in the base elicited prior is 50%. In other words, \( \varepsilon = 0.5 \) means that we elicit the \( \pi_0 \) prior but feel we could be as much as 50% off (in terms of implied probability sets).

9We chose: \( \theta_0 = 0, b_0 = 0 \) and \( \tau = 1 \).
model is usually estimated using GMM (see Arellano and Bond (1991); Blundell and Bond (1998), amongst others). It could also be estimated using the quasi-maximum likelihood (QML) estimator (see Bhargava and Sargan (1983), Kripfganz (2016), Bun et al. (2017), Moral-Benito et al. (2019)). Thus we compare our Bayesian two stage estimator with the Arellano-Bond GMM and the QML estimators.

Table 1 reports the results of fitting the Bayesian two stage model with block resampling boot-

strap (2S bootstrap)\(^\text{11}\) along with those from the GMM and QMLE, each in a separate panel respectively for \((N = 100, T = 10)\) and \((N = 200, T = 30)\). The true parameter values appear in the first row of the Table. The last column reports the computation time in seconds.\(^\text{12}\) Note that the computation time increases significantly as we move from a small sample to a larger one (the QMLE being the fastest).

The first noteworthy feature of the Table is that all the estimators yield parameter estimates, standard errors\(^\text{13}\) and RMSEs that are very close. For the coefficient of the lagged dependent variable, \(\rho\), the RMSE is the lowest for the 2S bootstrap when \(N = 100\) and \(T = 10\), but this RMSE is the lowest for the QMLE when \(N = 200\) and \(T = 30\), although the differences are small. GMM yields higher RMSEs for all coefficients. For the \(\beta\) coefficients, results are mixed in terms of RMSE for \(N = 100\) and \(T = 10\), but QMLE is the best for \(N = 200\) and \(T = 30\) with still small differences. The 2S bootstrap has better RMSEs than the frequentist estimators (GMM and QMLE) for the residual disturbances \((\sigma^u_\theta)\) and the random effects \((\sigma^\mu_\theta)\). Table I confirms that the base prior is not consistent with the data since \(\lambda_{\theta, g_0}\) is close to zero. The ML-II posterior density of \(\theta\) is close to the posterior \(\hat{q}^*(\theta|g_0)\) and to the empirical Bayes estimator \(\hat{\theta}_{EB}(b|g_0)\). In contrast, \(\hat{\lambda}_\mu\) is close to 0.5 so the Bayesian estimator \(b_\theta(\theta|g_0)\) under the base prior \(h_0\) and the empirical Bayes estimator \(\hat{b}_{EB}(\theta|h_0)\) each contribute similarly to the random effects \(b_i(\equiv \mu_i)\).

Table B.2 in the supplementary material gives the results when the coefficient \(\rho\) of the lagged dependent variable is increased from 0.75 to 0.98 (close to the unit root) for \(N = 100\) and \(T = 10\). The GMM estimator performs the worst as compared to the two other estimators. Even with a coefficient \(\rho\) very close to the unit root, the 95\% Highest Posterior Density Interval (HPDI) of the Bayesian estimator confirm the stationarity of the \(AR(1)\) process. It does not therefore seem necessary to impose a stationarity constraint on the prior distribution of \(\rho\). QMLE has the lowest RMSE (although the differences are small) except for \(\sigma^u_\theta\) and \(\sigma^\mu_\theta\) where 2S bootstrap is the best as reported in Table 1.

\(^{10}\)We use our own R codes for our Bayesian estimator, the R package “plm” for the Arellano-Bond GMM estimator and the “xtglsdpdm1” Stata package. We use the same DGP set under R and Stata environments to compare the three methods. We thank Jean-Michel Etienne for his help and support with the full-blown programming language Mata of Stata.

\(^{11}\)Recall that we use only \(BR = 20\) individual block bootstrap samples. Fortunately, the results are very robust to the value of \(BR\). For instance, increasing \(BR\) from 20 to 200 in the random effects world increases the computation time tenfold but yields practically the same results.

\(^{12}\)The simulations were conducted using R version 3.3.2 on a MacBook Pro, 2.8 GHz core i7 with 16Go 1600 MGz DDR3 ram.

\(^{13}\)Strictly speaking, we should mention “posterior means” and “posterior standard errors” whenever we refer to Bayesian estimates and “coefficients” and “standard errors” when discussing frequentist ones. For the sake of brevity, we will use “coefficients” and “standard errors” in both cases.
3.1.2. The Chamberlain-type fixed effects world

For the Chamberlain (1982)-type specification, the individual effects are given by \( \mu = X\beta + \omega \), where \( X \) is a \((N \times (T-1)K_x)\) matrix with \( X_1 = (X_{i1}, \ldots, X_{iT}) \) and \( \beta = (\pi_{i1}, \ldots, \pi_{iT})\) is a \(((T-1)K_x \times 1)\) vector. Here \( \pi_t \) is a \((K_x \times 1)\) vector of parameters to be estimated. The model can be rewritten as: \( y = Z\theta + Z_\mu X\beta + Z_\omega \omega + u \). We concatenate \([Z, Z_\mu X]\) into a single matrix of observables \( Z^* \) and let \( Wb = Z_\mu \omega \).

For the Chamberlain world, we compare the QML estimator to our Bayesian estimator. These are based on the transformed model: \( y_{it} = \rho y_{i,t-1} + x_{1,i,t} \beta_1 + x_{1,2,i,t} \beta_2 + \sum_{t=2}^{T} x_{2,i,t} \pi_t + \omega_i + u_{it} \) or \( y = Z^* \theta^* + Wb + u = Z^* \theta^* + Z_\mu \omega + u \) where \( Z^* = [y_{-1}, x_{1,1}, x_{1,2}, x_{2,2}] \), \( W = Z_\mu \) and \( b = \omega \).

Table 2 once again shows that the results of the 2S bootstrap are very close to those of the QML estimator. 2S bootstrap has the lowest RMSE for \( N = 100 \) and \( T = 10 \) except for \( \sigma_y^2 \). QMLE has the lowest RMSEs for all the parameters when \( N = 200 \) and \( T = 30 \). Table B.3 in the supplementary appendix gives the estimates of the \( \pi_t \) coefficients. The RMSEs are lower for QMLE than 2S bootstrap, although again the differences are very small. Table B.4 in the supplementary material report the results for \( N = 200 \) and \( T = 30 \).

3.1.3. The Hausman-Taylor world

The static Hausman-Taylor model (henceforth HT, see Hausman and Taylor (1981)) posits that \( y = X\beta + V\eta + Z_\mu \mu + u \), where \( V \) is a vector of time-invariant variables, and that subsets of \( X \) (e.g., \( X_{2,i}^* \)) and \( V \) (e.g., \( V_{2i}^* \)) may be correlated with the individual effects \( \mu \), but leaves the correlations unspecified. Hausman and Taylor (1981) proposed a two-step IV estimator.

For our dynamic general model (8) and for equation (22): \( y = Z\theta + Wb + u = \rho y_{i,t-1} + X\beta + V\eta + Z_\mu \mu + u \), we assume that \((\bar{X}_{2,i}^*, V_{2i}')\) are jointly normally distributed:

\[
\begin{pmatrix}
\mu_i \\
\bar{X}_{2,i}'
\end{pmatrix} 
\sim N \begin{pmatrix}
0 \\
E_{X'X'} \text{ and } E_{V'V'}
\end{pmatrix},
\]

where \( \bar{X}_{2,i}' \) is the individual mean of \( X_{2,i}^* \). The conditional distribution of \( \mu_i \mid \bar{X}_{2,i}', V_{2i}' \) is given by:

\[
\mu_i \mid \bar{X}_{2,i}', V_{2i}' \sim N \left( \begin{pmatrix} \Sigma_{11} & \Sigma_{12} \\ \Sigma_{21} & \Sigma_{22} \end{pmatrix}^{-1} \begin{pmatrix} X_{2,i} - E_{X_{2,i}}' \\ V_{2i}' - E_{V_{2i}}' \end{pmatrix} , \Sigma_{11} - \Sigma_{12} \Sigma_{22}^{-1} \Sigma_{21} \right).
\]

Since we do not know the elements of the variance-covariance matrix \( \Sigma_{js} \), we can write:

\[
\mu_i = \left( \bar{X}_{2,i}' - E_{X_{2,i}}' \right) \theta_X + \left( V_{2i}' - E_{V_{2i}}' \right) \theta_V + \omega_i,
\]

where \( \omega_i \sim N \left( 0, \Sigma_{11} - \Sigma_{12} \Sigma_{22}^{-1} \Sigma_{21} \right) \) is uncorrelated with \( u_{it} \), and where \( \theta_X \) and \( \theta_V \) are vectors of parameters to be estimated. In order to identify the coefficient vector of \( V_{2i}' \) and to avoid possible collinearity problems, we assume that the individual effects are given by:

\[
\mu_i = \left( \bar{X}_{2,i}' - E_{X_{2,i}}' \right) \theta_X + \left( V_{2i}' - E_{V_{2i}}' \right) f \left( \left( \bar{X}_{2,i}' - E_{X_{2,i}}' \right) \odot (V_{2i}' - E_{V_{2i}}') \right) \theta_V + \omega_i,
\]

(24)

where \( \odot \) is the Hadamard product and \( f \left( \left( \bar{X}_{2,i}' - E_{X_{2,i}}' \right) \odot (V_{2i}' - E_{V_{2i}}') \right) \) can be a nonlinear function of \( \left( \bar{X}_{2,i}' - E_{X_{2,i}}' \right) \odot (V_{2i}' - E_{V_{2i}}') \). The first term on the right-hand side of equation (24) corresponds
to the Mundlak (1978) transformation while the middle term captures the correlation between $V_{z_1}$ and $\mu_i$. The individual effects, $\mu_i$, are a function of $PX$ and $(f[PX \odot V])$, i.e., a function of the column-by-column Hadamard product of $PX$ and $V$ where $P = (I_N \otimes J_{T-1})/(T-1)$ is the between transformation. We can once again concatenate $[y_{t-1}, X, PX, f[PX \odot V]]$ into a single matrix of observables $Z^*$ and let $Wb \equiv Z_{2sT}$.

For our model (22), $y_{it} = \rho y_{i,t-1} + x_{i,t-1,1} \beta_{1,1} + x_{i,t,1,2} \beta_{1,2} + x_{2,i,t} \beta_2 + V_1, \eta_1 + V_2, \eta_2 + \mu_i + u_{it}$ or $y = \rho y_{t-1} + X_1 \beta_1 + X_2 \beta_2 + V_1, \eta_1 + V_2, \eta_2 + Z_{it} \mu_i + u_{it}$. Then, we assume that

$$
\mu_i = (\bar{\sigma}_{2,T} - E_{\bar{\sigma}_{2,T}}) \theta_X + f ((\bar{\sigma}_{2,T} - E_{\bar{\sigma}_{2,T}}) \odot (V_{2i} - E_{V_{2i}})) \theta_V + \omega_i.
$$

We propose adopting the following strategy: If the correlation between $\mu_i$ and $V_{2i}$ is quite large ($> 0.2$), use $f[.] = (\bar{\sigma}_{2,T} - E_{\bar{\sigma}_{2,T}})^s \odot (V_{2i} - E_{V_{2i}})$ with $s = 1$. If the correlation is weak, set $s = 2$. In real-world applications, we do not know the correlation between $\mu_i$ and $V_{2i}$, a priori. We can use a proxy of $\mu_i$ defined by the OLS estimation of $\mu$: $\hat{\mu} = (Z^T_{2i} Z_{2i})^{-1} Z^T_{2i} \hat{\gamma}$ where $\hat{\gamma}$ are the fitted values of the pooling regression $y = \rho y_{t-1} + X_1 \beta_1 + X_2 \beta_2 + V_1, \eta_1 + V_2, \eta_2 + \gamma_i$. Then, we compute the correlation between $\hat{\mu}$ and $V_2$. In our simulation study, it turns out the correlations between $\mu_i$ and $V_2$ are large: 0.65. Hence, we choose $s = 1$. In this specification, $Z = [y_{t-1}, x_{t,1,1}, x_{t,1,2}, x_{t,2}, V_1, V_2, P_{x,t}, f[PX \odot V]]$, $W = Z_0$, and $b = \infty$.

Our 2S bootstrap estimation method is compared with the two-stage quasi-maximum likelihood sequential approach proposed by Kripfganz and Schwarz (2019). In the first stage, they estimate the coefficients of the time-varying regressors without relying on coefficient estimates for the time-invariant regressors using the quasi-maximum likelihood (QML) estimator of Hsiao et al. (2002) with the “xtdpdqml” Stata command. Subsequently, they regress the first-stage residuals on the time-invariant regressors. They achieve identification by using instrumental variables in the spirit of Hausman and Taylor (1981), and they adjust the second-stage standard errors to account for the first-stage estimation error.14 They have proposed a new “xtseqreg” Stata command which implements the standard error correction for two-stage dynamic linear panel data models.15

Table 3 compares results of the 2S bootstrap estimator to those of the two-stage QML sequential approach. Once again, the estimates are very close to one another. The RMSE is smaller for the two-stage QML than for the 2S bootstrap for all the parameters for $N = 200$ and $T = 10$. On the other hand, the 2S bootstrap has a lower RMSE for $\eta_2$ (for $N = 100$ and $T = 10$) and $(N = 100$ and $T = 30)$. This is true despite the fact that the 2S bootstrap estimator yields a slightly upward biased estimate of $\eta_2$, the coefficient associated with the time-invariant variable $Z_{it}$, which is itself correlated with $\mu_i$. This bias decreases as $T$ increases (from 16% for $T = 10$ to 4.7% for $T = 30$). Interestingly, the standard errors of that same coefficient $\eta_2$ are smaller when using the Bayesian estimator as compared to the two-stage QMLE, and especially when $T$ is larger. Even with a slight bias, the 95% confidence intervals of the Bayesian estimator are narrower and entirely nested within those obtained with the two-stage QML sequential approach. We also reached the same conclusion

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14For the following specification: $y_{it} = \rho y_{i,t-1} + x_{i,t} \beta + V_{i,t} \eta + \mu_i + u_{it}$, the first stage model is $y_{it} = \rho y_{i,t-1} + x_{i,t} \beta + \epsilon_{it}$, where $\epsilon_{it} = \kappa_i - \gamma + \mu_i$ and $\kappa_i = V_{i,t} \eta + \mu_i$ and is estimated in first differences. In the second stage, Kripfganz and Schwarz (2019) estimate the coefficients $\gamma$ based on the level relationship: $y_{it} - \rho y_{i,t-1} - x_{i,t} \beta = V_{i,t} \eta + \gamma_i$, where $\gamma_i = \mu_i + \gamma_i + (\beta - \gamma) y_{i,t-1} - x_{i,t} \beta - \beta$ and compute proper standard errors with an analytical correction term.

15Following Kripfganz and Schwarz (2019), we use successively these two Stata commands (“xtdpdqml” and “xtseqreg”). Unfortunately, these Stata commands do not give the residual variance of specific effects $\sigma_u^2$ but only $\sigma_u^2$. 

---
in a static model (see Baltagi et al. (2018)). Finally, note that the 2S bootstrap estimator yields slightly biased estimates of \( \sigma^2 \) but this bias decreases rapidly as the time span is increased (from 17.5% for \( T = 10 \) to 2.5% for \( T = 30 \)).

3.1.4. The dynamic homogeneous panel data world with common trends

The dynamic homogeneous panel data world with common trends is defined as:

\[
y_{it} = \rho y_{i,t-1} + x_{it} \beta_1 + x_{i,t-1} \beta_2 + f_t^\prime \gamma_i + u_{it}
\]

Since the \( m \) common trends, \( f_t \), are known, we can rewrite the general dynamic model (8) as follows:

\[
y = Z \theta + Wb + u = Z \theta + F \Gamma + u
\]

where \( u \sim N(0, \Sigma) \), \( \Sigma = \tau^{-1}I_N \). The \((N(T-1) \times Nm)\) matrix \( F \) of the \( m \) common trends is a blockdiagonal matrix where each \((T-1 \times m)\) sub block \( f \) is replicated \( N \) times and \( \Gamma \) is the \((Nm \times 1)\) individual varying coefficients vector:

\[
F = I_N \otimes f \text{ with } f = \begin{pmatrix} f_{21} & \ldots & f_{2m} \\ \ldots & \ldots & \ldots \\ f_{T1} & \ldots & f_{Tm} \end{pmatrix} \text{ and } \Gamma = vec\begin{pmatrix} \gamma_{11} & \gamma_{21} & \ldots & \gamma_{N1} \\ \gamma_{12} & \gamma_{22} & \ldots & \gamma_{N2} \\ \ldots & \ldots & \ldots & \ldots \\ \gamma_{1m} & \gamma_{2m} & \ldots & \gamma_{Nm} \end{pmatrix}
\]

This model is usually estimated using the common correlated effects pooled estimator (CCEP) (see Pesaran (2006) and Chudik and Pesaran (2015a,b)). It can also be estimated using the quasi-maximum likelihood (QML) estimator. We compare our 2S bootstrap estimator with the CCEP estimator. We use our own R codes for our Bayesian estimator and the “xtdece2” Stata package for the CCEP estimator. We use the same DGP set under R and Stata environments to compare the two methods.

Table 4 shows that the results of the 2S bootstrap estimator are close to those of the CCEP estimator. The results on RMSEs are mixed. 2S bootstrap gives a lower RMSE for \( \rho \) than CCEP for \( N = 100 \) and \( T = 30 \), i.e. 0.002 compared to 0.007, but this difference is reduced for \( N = 50 \) and \( T = 50 \), i.e. 0.0058 compared to 0.0066. CCEP gives a lower RMSE for \( \beta_1 \). For \( \beta_2 \), it depends on the sample sizes. Finally, 2S bootstrap gives a lower RMSE for \( \sigma^2 \). The computation time is a bit longer with our estimator given the bootstrap procedure. However, all estimators yield essentially the same parameter estimates, no matter what sample size.

3.1.5. The dynamic homogeneous panel data world with correlated common effects

Again, this model is usually estimated using the common correlated effects pooled estimator (CCEP) (see Pesaran (2006); Chudik and Pesaran (2015a,b)) or with the principal components estimators using quasi-maximum likelihood (QML) estimator (see Bai (2009) or Song (2013)). Since the \( m \) common correlated effects, \( f_t \), are unknown, we need to rewrite the general dynamic

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**Note:**

16 We use our own R codes for our Bayesian estimator and the “xtdece2” Stata package for the CCEP estimator. We use the same DGP set under R and Stata environments to compare the two methods.
model (8) as follows:

\[ y = Z\theta + Wh + u = Z\theta + FT + u \]

with \( Z'_{it} = [y_{it-1}, X'_{it}] \), \( \theta' = [\rho, \beta]' \) and \( X'_{it} = [x_{it}, x_{it-1}] \),

where the \((N(T - 1) \times Nm)\) matrix \( F \) of the \( m \) unobserved factors is still a block diagonal matrix where each \(((T - 1) \times m)\) sub block \( f \) is replicated \( N \) times but \( f \) should be approximated by known variables. Similar to the Hausman-Taylor case (see eq(24)), we can approximate the \(((T - 1) \times m)\) \( f \) matrix with a \(((T - 1) \times K_i)\) \( f^* \) matrix of the within time transformation\(^1\) of \( Z_{it} \):

\[ f^* = \begin{pmatrix} f^* \vdots f^* \\ f^* \end{pmatrix} \]

where \( f^* = [(\overline{y}_{-1,i} - \overline{y}_{-1}), (\overline{x}_i - \overline{x}), (\overline{x}_{-1,i} - \overline{x}_{-1})] \)

with \( \overline{x}_i = (1/N) \sum_{i=1}^{N} x_{it}, \overline{y} = (1/NT) \sum_{t=1}^{T} \sum_{i=2}^{N} x_{it}, \)

or as Chudik and Pesaran (2015a) by the time means of the dependent and explanatory variables:

\[ f^*_t = [\overline{y}_{-1,i}, \overline{x}_i, \overline{x}_{-1,i}] \].\(^1\)

We follow the method of Chudik and Pesaran (2015a,b) by introducing the time means of the dependent and explanatory variables instead of introducing only the within time transformation of the explanatory variables \( Z'_{it} \). Then, the product \( FT \) is approximated with the product \( F^*\Gamma^* \) where the factor loadings \( \Gamma^* \) is a \((NK_i \times 1)\) vector and \( F^* \) is a \((N(T - 1) \times NK_i)\) matrix of the time means of \( Y \) and \( Z \).

Table 5 shows that the results of the 2S bootstrap are very close to those of the dynamic CCEP estimator. However, the RMSE is smaller for CCEP than 2S bootstrap, but not by much for most parameters.

3.1.6. The dynamic heterogeneous panel data world with correlated common effects

The dynamic heterogeneous panel data world with common factors is defined as:

\[ y_{it} = \rho y_{i,t-1} + x_{it}\beta_{1i} + x_{i,t-1}\beta_{2i} + f^*_{i}\gamma_i + u_{it} = Z'_{it}\theta_i + f^*_{i}\gamma_i + u_{it} \]

where \( Z'_{it} = [y_{it-1}, X'_{it}] \), \( \theta'_i = [\rho_i, \beta_{1i}', \beta_{2i}' \) and \( X'_{it} = [x_{it}, x_{it-1}] \). This model is usually estimated using the common correlated effects mean group estimator (CCEMG) (see Pesaran (2006) and Chudik and Pesaran (2015a,b)). It could also be estimated using the quasi-maximum likelihood (QML) estimator. So we compare the mean coefficients \( \hat{\theta} = (1/N) \sum_{i=1}^{N} \hat{\theta}_i \) of our 2S bootstrap estimator with the CCEMG estimator.\(^1\)

While the bottom panel of Table 6 gives insights on the distribution of \( \rho_i \) and \( \beta_{1i} \) for different sample sizes, the top panel of Table 6 gives the estimated values of the mean coefficients \( \overline{\rho}, \overline{\beta_1}, \) the estimated values of \( \beta_2 \) and \( \sigma_u^2 \), their standard deviations and their RMSE’s. Table 6 shows that

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\(^1\)i.e., the demeaned time means.

\(^1\)The dynamic CCEP estimator is defined as: \( y_{it} = \rho y_{i,t-1} + x_{it}\beta_1 + x_{i,t-1}\beta_2 + \sum_{j=0}^{P_T} f^*_{i-j}\gamma_{i,j} + u_{it} \) where \( P_T = T^{1/3} \) (see Chudik and Pesaran (2015b) pp. 26). Then, \( P_T \approx 3 \) when \( T = 30 \) or \( T = 50 \). In the simulations, we use \( P_T = 0 \).

\(^1\)We use our own R codes for our Bayesian estimator and the “xttdc2” Stata package for the CCEMG estimator. We use the same DGP set under R and Stata environments to compare the two methods.
the results of the 2S bootstrap estimator are close to those of the CCEMG estimator. The RMSEs results are mixed. 2S bootstrap gives a smaller RMSE for \( \rho \) than CCEMG, but CCEMG gives a smaller RMSE for \( \beta \) and \( \sigma^2 \). The results for \( \beta \) depend on the sample size. For \( N = 50 \), the bias is (7.64\% for \( N = 100 \), \( T = 30 \)), (resp. 16.47\% for \( N = 50 \), \( T = 50 \)) for the 2S bootstrap estimator as compared to those of the CCEMG estimator (4.37\%, resp.10.21\%). For the residuals' variance \( \sigma^2 \), the bias increases with the time dimension for both estimators. However, all the estimators yield roughly the same parameter estimates. Computation time is a bit longer with our estimator given the bootstrap procedure.

3.1.7. Sensitivity to \( \varepsilon \)-contamination values

Tables B.5 and B.6 (in the supplementary material) investigate the sensitivity of the 2S bootstrap estimator for the random effects world and for the heterogeneous panel data world with correlated common effects\(^{20}\) with respect to \( \varepsilon \), the contamination part of the prior distributions, which varies between 0 and 90\%. As shown in Table B.5 for the random effects world when \( N = 100 \) and \( T = 10 \), all the parameter estimates are insensitive to \( \varepsilon \). The only noteworthy change concerns the estimated value of \( \lambda_\mu(\equiv \lambda_{b,h_{\epsilon}}) \). It more or less corresponds to \( (1 - \varepsilon) \). This particular relation may occur whenever \( \tilde{h}/(\tilde{h} + 1) = h_0/(h_0 + 1) \) and \( R_{b,h}^2/\left(1 - R_{b,h}^2\right) = R_{b,\epsilon}^2/\left(1 - R_{b,\epsilon}^2\right) \) (see the definition of \( \lambda_{b,h_{\epsilon}} \) in section 2.3.2). The observed stability of the coefficients estimates stems from the fact that the base prior is not consistent with the data as the weight \( \lambda \) is implicitly assumed error-free. This is a strong assumption. This time, results are not strictly similar to those of \( \varepsilon = 0 \) constrains the model to rely exclusively on a base elicited prior which is implicitly assumed error-free. This is a strong assumption. This time, results are not strictly similar to those of \( \varepsilon = 0 \) but they are close to the true values except for \( \sigma^2 \) which has a fairly large upward bias (11.4\%) as well as a large RMSE.\(^{21}\)

Table B.6 shows similar results. All the parameter estimates are insensitive to \( \varepsilon \) (\( \varepsilon \neq 0 \)) for the heterogeneous panel data world with correlated common effects when \( N = 100 \) and \( T = 30 \). The only changes concern the estimated values of \( \lambda_{\theta,g_0} \) and \( \lambda_\mu(\equiv \lambda_{b,h_{\epsilon}}) \). While \( \lambda_\mu(\equiv \lambda_{b,h_{\epsilon}}) \) changes inversely to \( \varepsilon \), \( \lambda_{\theta,g_0} \) has the shape of an inverted J as \( \varepsilon \) increases. As for the random effects world, when \( \varepsilon = 0 \), the results are not strictly similar to those of \( \varepsilon = 0 \) but they are close to the true values except for \( \sigma^2 \) which has also a fairly large upward bias (14.5\%) as well as large standard error and RMSE. Whatever the world tested, results are insensitive to the exact value of \( \varepsilon \). This stems from the fact that the 2S bootstrap estimator is data driven and implicitly adjusts the weights to the different values of \( \varepsilon \)-contamination. This may by why, even though the choice of \( \varepsilon = 0.5 \) is somewhat arbitrary, the adjustment compensates for it not being optimal (see Berger (1985)).

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\(^{20}\)This exercise could be conducted for the other worlds as Chamberlain-type fixed effects or Hausman-Taylor world but we report the results for only two worlds for the sake of brevity.

\(^{21}\)From a theoretical point of view, and under the null, \( H_0 : \varepsilon = 0 \), it follows that the weights \( \lambda_{\theta,g_0} = 1 \) and \( \lambda_{b,h_{\epsilon}} = 1 \) so that the restricted ML-II estimator of \( \theta \) is given by \( \hat{\theta}_{\text{restrict}} = \theta_*(b \mid g_0) \). Under \( H_1 : \varepsilon \neq 0 \) the unrestricted estimator is \( \hat{\theta}_{\text{unrestrict}}(\equiv \hat{\theta}_{\text{ML-II}}) = \lambda_{\theta,g_0}\theta_*(b \mid g_0) + \left(1 - \lambda_{\theta,g_0}\right)\theta_{\text{EB}}(b \mid g_0) \). The restricted ML-II estimator \( \theta_*(b \mid g_0) \) is the Bayes estimator under the base prior \( g_0 \).
3.1.8. Departure from normality

Tables B.7 and B.8 (in the supplementary material) investigate the robustness of the estimators to a non-normal framework for the random effects world and for the heterogeneous panel data world with correlated common effects. The remainder disturbances, \( u_{it} \), are now assumed to follow a right-skewed \( t \)-distribution with mean = 0, degrees of freedom \( \nu = 3 \), and skewing parameter \( \gamma = 2 \) (see Fernández and Steel (1998), Baltagi et al. (2018)). Our 2S bootstrap estimator behaves pretty much like the GMM and the QMLE for the random effects world when \( N = 100 \) and \( T = 10 \) (see Table B.7). There is, however, a slight downward bias for the \( \rho \) coefficient (−4.7%) and a slight upward bias for \( \beta_{11} \) and \( \beta_2 \) (7%). But these biases are small. Compared to the GMM estimator, our 2S bootstrap estimator provides better estimates of \( \sigma_u^2 \) and \( \sigma^2 \) but it is the QML estimator that gives the estimates closest to the true values. Another interesting result concerns the standard errors and RMSEs of all the estimators. The presence of a right-skewed \( t \)-distribution greatly increases these values especially for \( \sigma_u^2 \).

Table B.8 investigates the robustness of the CCEMG and 2S bootstrap estimators to the right-skewed \( t \)-distribution for the heterogeneous panel data world with correlated common effects when \( N = 100 \) and \( T = 30 \). There are slight downward biases for the \( \overline{\rho} \) mean coefficient, with that of CCEMG being larger than that of 2S bootstrap (−16% vs −7.6%) as well as slight upward bias for \( \sigma_u^2 \), that of CCEMG being larger than that of 2S bootstrap (5.6% vs 0.5%). However, for the \( \overline{\beta}_1 \) mean coefficient, it is the 2S bootstrap estimator which has a slightly larger bias (17.2% vs 7.5%). Finally, it can be noted that the RMSE of \( \sigma_u^2 \) is larger for the CCEMG estimator than for the 2S bootstrap estimator.

4. Conclusion

To our knowledge, our paper is the first to analyze the dynamic linear panel data model using an \( \epsilon \)-contamination approach within a two-stage hierarchical approach. The main benefit of this approach is its ability to extract more information from the data than the classical Bayes estimator with a single base prior. In addition, we show that our approach encompasses a variety of classical or frequentist specifications. We estimate the Type-II maximum likelihood (ML-II) posterior distribution of the slope coefficients and the individual effects using a two-step procedure. The posterior distribution is a convex combination of the conditional posterior densities derived from the elicited prior and the \( \epsilon \)-contaminated prior. Thus if the base prior is consistent with the data, more weight is given to the conditional posterior density derived from the former. Otherwise, more weight is given to the latter.

The finite sample performance of the two-stage hierarchical models is investigated using extensive Monte Carlo experiments. The experimental design includes a random effects world, a Chamberlain type fixed effects world, a Hausman-Taylor-type world and worlds with homogeneous/heterogeneous slopes and cross-sectional dependence. The simulation results underscore the relatively good performance of the two-stage hierarchy estimator, irrespective of the data generating process considered. The biases and the RMSEs are close and sometimes smaller than those of

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22 The Skewed \( t \) distribution with \( \nu \) degrees of freedom and skewing parameter \( \gamma \) has the following density:

\[
pdf(x) = \frac{2}{\gamma + x} f(z) \text{ where } z = \gamma x \text{ if } x < 0 \text{ or } z = x/\gamma \text{ if } x \geq 0
\]

where \( f(.) \) is the density of the \( t \) distribution with \( \nu \) degrees of freedom.
the conventional (classical) estimators. We also investigate the sensitivity of the estimators to the contamination part of the prior distribution. It turns out that parameter estimates are relatively stable. Finally, we investigate the robustness of the estimators when the remainder disturbances are assumed to follow a right-skewed $t$-distribution. Compared to classical estimators, our robust estimators globally behave well in terms of precision and bias.

The robust Bayesian approach we propose is arguably a useful all-in-one panel data estimator. Because it embeds a variety of estimators, it can be used straightforwardly to estimate dynamic panel data models under many alternative stochastic specifications. Unlike classical estimators, there is no need to have a custom estimator for each possible DGP. Furthermore, it allows one to circumvent the difficulties faced by analysts who are oftentimes constrained to use those estimators that are readily available in standard software suites.

We reckon that our estimator contributes only marginally to those already available in the literature. Our main contribution is to propose an estimator that allows the analysts to focus on the stochastic specification of their model rather than finding the software best suited to their needs. This is because our estimator is easily amenable to many specifications in addition to those already presented in this paper. These include models with individual and time random effects in unknown common factors, spatial structures (autoregressive spatial), etc. We leave these for future work.
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<th>β₁₂</th>
<th>β₂</th>
<th>σ²ᵤ</th>
<th>σ²ₚ</th>
<th>λ₀</th>
<th>λₘ</th>
<th>Computation Time (secs.)</th>
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2S boot: two stage with individual block resampling bootstrap.
GMM: Arellano-Bond GMM estimation.
QMLE: quasi-maximum likelihood estimation.
Table 2: Dynamic Chamberlin-type Fixed Effects World ($\varepsilon = 0.5$)

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<th>$\beta_{11}$</th>
<th>$\beta_{12}$</th>
<th>$\beta_2$</th>
<th>$\sigma_u^2$</th>
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<th>$\lambda_0$</th>
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<td>1.0132</td>
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2S boot: two stage with individual block resampling bootstrap.

The parameters $\pi_t$ are omitted from the table, see Table B.3 in appendix B in the supplementary material.

QMLE: quasi-maximum likelihood estimation.

The parameters $\pi_t$ are omitted from the table, see Table B.4 in appendix B in the supplementary material.

(*) When $T = 30$, we restrict the exercise to only 500 replications, not because of our estimator under R but because of the size of the simulated database and its reading under Stata. Indeed, for 1,000 replications and $T = 30$, one must read, under Stata, $(1 + 4 + 30) \times 1,000 = 35,000$ variables of size $(NT, 1)$! Even with a 64-bit computer and Stata (MP and S), only 32,767 variables can be read. On the other hand, with our code under R, there is no limitation of that order.
Table 3: Dynamic Hausman-Taylor World
\( \rho = 0.8, \varepsilon = 0.5, \text{replications}=1,000 \)

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<th>( \beta_2 )</th>
<th>( \eta_1 )</th>
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<th>( \sigma_u^2 )</th>
<th>( \sigma_\mu^2 )</th>
<th>( \lambda_\theta )</th>
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<td>n.a</td>
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</tr>
</tbody>
</table>

2S boot: two stage with individual block resampling bootstrap.

two-stage QML: two-stage quasi-maximum likelihood sequential approach with non available (n.a) estimate of \( \sigma_\mu^2 \).
Table 4: Dynamic homogeneous panel data model with common trends
$\varepsilon = 0.5$, Replications=1,000

<table>
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<th>$\beta_2$</th>
<th>$\sigma_u^2$</th>
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<th>$\lambda_\mu$</th>
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2S boot: two stage with individual block resampling bootstrap.
CCEP: Common Correlated Effects Pooled estimator.
Table 5: Dynamic homogeneous panel data model with common correlated effects
\( \varepsilon = 0.5, \) Replications=1,000

<table>
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<tr>
<th></th>
<th>( \rho )</th>
<th>( \beta_1 )</th>
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<th>( \sigma_u^2 )</th>
<th>( \lambda_{\theta} )</th>
<th>( \lambda_{\mu} )</th>
<th>Computation Time (secs.)</th>
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2S boot: two stage with individual block resampling bootstrap.
CCEP: Common Correlated Effects Pooled estimator.
Table 6: Dynamic heterogeneous panel data model with common correlated effects

\( \varepsilon = 0.5, \text{Repetitions}=1,000 \)

| Mean coefficients: \( \bar{\rho} = \frac{1}{N} \sum_{i=1}^{N} \rho_i \) and \( \bar{\beta}_1 = \frac{1}{N} \sum_{i=1}^{N} \beta_{1i} \). |
|---|---|---|---|---|---|---|---|
| true | \( \bar{\rho} \) | \( \bar{\beta}_1 \) | \( \bar{\beta}_2 \) | \( \sigma^2_{\nu} \) | \( \lambda_\theta \) | \( \lambda_\mu \) | Computation Time (secs.) |
| \( N = 100 \) | 2S boot coef | 0.7347 | 0.8071 | 1.0759 | 1.0536 | 0.0011 | 0.3925 | 4695.86 |
| | se | 0.0194 | 0.0671 | 0.0531 | 0.0392 | | | |
| | rmse | 0.0248 | 0.0882 | 0.0926 | 0.0664 | | | |
| | CCEMG coef | 0.7205 | 0.7826 | 1.0856 | 1.0409 | | | 1618.05 |
| | se | 0.0221 | 0.0554 | 0.0453 | 0.0353 | | | |
| | rmse | 0.0369 | 0.0600 | 0.0968 | 0.0540 | | | |
| \( N = 50 \) | 2S boot coef | 0.7474 | 0.8739 | 1.0666 | 1.0838 | 0.0010 | 0.4407 | 2492.75 |
| | se | 0.0217 | 0.0973 | 0.0643 | 0.0474 | | | |
| | rmse | 0.0219 | 0.1572 | 0.0925 | 0.0963 | | | |
| | CCEMG coef | 0.7423 | 0.8269 | 1.0654 | 1.1450 | | | 1145.23 |
| | se | 0.0233 | 0.0848 | 0.0619 | 0.2180 | | | |
| | rmse | 0.0245 | 0.1159 | 0.0900 | 0.2618 | | | |

2S boot: two stage with individual block resampling bootstrap.
CCEMG: Common Correlated Effects Mean Group estimator.

\( N = 100, T = 30 \)

\( N = 50, T = 50 \)

<table>
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<th>( \rho_i )</th>
<th>( \beta_{1i} )</th>
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<td>max</td>
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Robust dynamic panel data models using $\varepsilon$-contamination

Supplementary material

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March 7, 2020

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A. Uniform distribution, derivation of the mean and variance of the ML-II posterior density of $\rho$ and some Monte Carlo results

A.1. Uniform distribution and derivation of the mean and variance of the ML-II posterior density of $\rho$

Following Singh and Chaturvedi (2012) (see also Shrivastava et al. (2019)), and for deriving the posterior density of $\rho$, given $(\beta, b)$, we write:

$$y^o = (y - X\beta - Wb) = \rho y_{-1} + u$$

the probability density function (pdf) of $y^o$, given the observables and the parameters, is:

$$p(y^o | y_{-1}, \rho, \tau) = \left(\frac{\tau}{2\pi}\right)^{\frac{N(T-1)}{2}} \exp \left(-\frac{\tau}{2}(y^o - \rho y_{-1})'(y^o - \rho y_{-1})\right)$$

Let $\hat{\rho}(\beta, b) = (y'_{-1}y_{-1})^{-1} y'_{-1} y^o = (\Lambda_y)^{-1} y'_{-1} y^o$, then following the derivation of (eq.16) in the technical appendix (pp 6-7) of Baltagi et al. (2018), we can write:

$$(y^o - \rho y_{-1})'(y^o - \rho y_{-1}) = \varphi(\beta, b) + \Lambda_y \{\rho - \hat{\rho}(\beta, b)\}^2$$

with

$$\varphi(\beta, b) = (y^o - \hat{\rho}(\beta, b)y_{-1})'(y^o - \hat{\rho}(\beta, b)y_{-1})$$

and

$$\hat{\rho}(\beta, b) = (y'_{-1}y_{-1})^{-1} y'_{-1} y^o = \Lambda_y^{-1} y'_{-1} y^o$$

then

$$p(y^o | y_{-1}, \rho, \tau) = \left(\frac{\tau}{2\pi}\right)^{\frac{N(T-1)}{2}} \exp \left(-\frac{\tau}{2} \left\{ \varphi(\beta, b) + \Lambda_y \{\rho - \hat{\rho}(\beta, b)\}^2 \right\} \right)$$

As the precision $\tau$ is assumed to have a vague prior $p(\tau) \propto \tau^{-1}$ and as $|\rho| < 1$ is assumed to be $U(-1, 1)$, the conditional posterior density of $\rho$, given $(\beta, b)$ is defined by:

$$\pi^*(\rho | \beta, b) = \int_0^\infty \int_{-1}^1 p(\rho)p(\tau) p(y^o | y_{-1}, \rho, \tau) d\tau d\rho$$

(A.1)

where

$$p(\rho) = \frac{1}{2} \text{ and } p(\tau) = \frac{1}{\tau}$$
So, the numerator of $\pi^* (\rho|\beta, b)$ can be written as:

$$
\int_0^\infty p(\rho) p(\tau) p(y^n|y_{-1}, \rho, \tau) \, d\tau = \int_0^\infty \left( \frac{1}{2} \right) \left( \frac{1}{\tau} \right) \left( \frac{\tau}{2\pi} \right)^{N(T-1)/2} \times \exp \left( -\frac{\tau}{2} \left\{ \phi(\beta, b) + \Lambda_y \{ \rho - \hat{\rho}(\beta, b) \}^2 \right\} \right) \, d\tau = \left( \frac{1}{2} \right) \left( \frac{1}{\tau} \right) \left( \frac{\tau}{2\pi} \right)^{N(T-1)/2} \times \int_0^\infty \exp \left( -\frac{\tau}{2} \left\{ \phi(\beta, b) + \Lambda_y \{ \rho - \hat{\rho}(\beta, b) \}^2 \right\} \right) \, d\tau
$$

As $\int_0^\infty \tau^{x-1} \exp \left( -\frac{\tau}{2} \right) \, d\tau = (2/\pi)^x \Gamma(x)$, then

$$
\int_0^\infty p(\rho) p(\tau) p(y^n|y_{-1}, \rho, \tau) \, d\tau = \Gamma \left( \frac{N(T-1)}{2} \right) \left( \frac{2}{\pi} \right)^{-\frac{N(T-1)}{2}} \left[ \phi(\beta, b) + \Lambda_y \{ \rho - \hat{\rho}(\beta, b) \}^2 \right]^{-\frac{N(T-1)}{2}}
$$

The denominator of the conditional posterior density of $\rho$ is

$$
\int_0^\infty \int_{-1}^1 p(\rho) p(\tau) p(y^n|y_{-1}, \rho, \tau) \, d\rho d\tau = \Gamma \left( \frac{N(T-1)}{2} \right) \left( \frac{2}{\pi} \right)^{-\frac{N(T-1)}{2}} \int_{-1}^1 \left[ \phi(\beta, b) + \Lambda_y \{ \rho - \hat{\rho}(\beta, b) \}^2 \right]^{-\frac{N(T-1)}{2}} \, d\rho
$$

then, the conditional posterior density of $\rho$, given $(\beta, b)$ is defined by:

$$
\pi^* (\rho|\beta, b) = \frac{\left[ \phi(\beta, b) + \Lambda_y \{ \rho - \hat{\rho}(\beta, b) \}^2 \right]^{-\frac{N(T-1)}{2}}}{\int_{-1}^1 \left[ \phi(\beta, b) + \Lambda_y \{ \rho - \hat{\rho}(\beta, b) \}^2 \right]^{-\frac{N(T-1)}{2}} \, d\rho}
$$

(A.2)
Let us derive the denominator of the previous expression

\[ A = \int_{-1}^{1} \left[ \varphi(\beta, b) + \Lambda_y \left\{ \rho - \hat{\rho}(\beta, b) \right\}^2 \right]^{\frac{N(T-1)}{2}} d\rho \]

\[ = [\varphi(\beta, b)]^{\frac{N(T-1)}{2}} \int_{-1}^{1} \left[ 1 + \Lambda_y \frac{\left\{ \rho - \hat{\rho}(\beta, b) \right\}^2}{\varphi(\beta, b)} \right]^{\frac{N(T-1)}{2}} d\rho \]

\[ = [\varphi(\beta, b)]^{\frac{N(T-1)}{2}} \sqrt{\frac{\Lambda_y}{\varphi(\beta, b)}} \left[ \int_{-\sqrt{\frac{\Lambda_y}{\varphi(\beta, b)}}(1-\hat{\rho}(\beta, b))}^{\sqrt{\frac{\Lambda_y}{\varphi(\beta, b)}}(1+\hat{\rho}(\beta, b))} \left( 1 + \frac{t^2}{1+t^2} \right) d\varphi \right]^{\frac{N(T-1)}{2}} \]

\[ = [\varphi(\beta, b)]^{\frac{N(T-1)}{2}} \sqrt{\frac{\Lambda_y}{\varphi(\beta, b)}} \left[ \int_{0}^{\sqrt{\frac{\Lambda_y}{\varphi(\beta, b)}}(1-\hat{\rho}(\beta, b))} \left( 1 + \frac{t^2}{1+t^2} \right) d\varphi + \int_{0}^{\sqrt{\frac{\Lambda_y}{\varphi(\beta, b)}}(1+\hat{\rho}(\beta, b))} \left( 1 + \frac{t^2}{1+t^2} \right) d\varphi \right]^{\frac{N(T-1)}{2}} \]

\[ = [\varphi(\beta, b)]^{\frac{N(T-1)}{2}} \sqrt{\frac{\Lambda_y}{\varphi(\beta, b)}} \left[ I_1 + I_2 \right] \]

Now taking the transformation \( \eta = \frac{t^2}{1+t^2} \), then

\[ \left( 1 + \frac{t^2}{1+t^2} \right) = \left( 1 - \eta \right)^{-1} \text{ and } dt = \frac{1}{2\eta^{\frac{3}{2}}} \left( 1 - \eta \right)^{-3/2} d\eta \]
and we obtain $I_1$ as

$$I_1 = \int_0^1 (1 + t^2) \frac{\varphi(t)}{\varphi(t)^2} \frac{\varphi(t)^{N(T-1)}}{\varphi(t)^2} \frac{d\varphi}{\varphi(t)}$$

where $\varphi = (1 + \bar{\rho}(\beta, b))^2$.

$$= \frac{1}{\sqrt{N(T-1)}} \int_0^1 (1 + t^2) \frac{\varphi(t)}{\varphi(t)^2} \frac{\varphi(t)^{N(T-1)-3}}{\varphi(t)^2} \frac{d\varphi}{\varphi(t)}$$

where $\varphi_1 = \frac{\Delta_\varphi(\beta, b)}{\varphi(\beta, b)} (1 + \bar{\rho}(\beta, b))^2$.

Using the Pfaff’s transformation:

$$\frac{\varphi(t)}{\varphi(t)^2} \frac{\varphi(t)^{N(T-1)-3}}{\varphi(t)^2} = \frac{1}{\sqrt{N(T-1)}} \int_0^1 (1 - \varphi(t)^{N(T-1)-3}) \frac{d\varphi}{\varphi(t)}$$

we obtain

$$= \frac{1}{\sqrt{N(T-1)}} \int_0^1 (1 - \varphi(t)^{N(T-1)-3}) \frac{d\varphi}{\varphi(t)}$$

Notice that

$$\frac{\varphi(t)}{\varphi(t)^2} \frac{\varphi(t)^{N(T-1)-3}}{\varphi(t)^2} = \frac{1}{\sqrt{N(T-1)}} \int_0^1 (1 - \varphi(t)^{N(T-1)-3}) \frac{d\varphi}{\varphi(t)}$$

Hence

$$I_1 = \frac{1}{\sqrt{N(T-1)}} \int_0^1 (1 + t^2) \frac{\varphi(t)}{\varphi(t)^2} \frac{\varphi(t)^{N(T-1)-3}}{\varphi(t)^2} \frac{d\varphi}{\varphi(t)}$$

Similarly we obtain

$$I_2 = \frac{1}{\sqrt{N(T-1)}} \int_0^1 (1 + t^2) \frac{\varphi(t)}{\varphi(t)^2} \frac{\varphi(t)^{N(T-1)-3}}{\varphi(t)^2} \frac{d\varphi}{\varphi(t)}$$

\[1\] The Euler integral formula is given by:

$$\int_0^1 (t^{a_2-1}) (1 - t^{a_3-a_2-1}) (1 - zt)^{-a_1} dt = \Gamma(a_2) \Gamma(a_2 - a_2) \Gamma(a_3) \times F_1 (a_1; a_2; a_3; z)$$

where $F_1 (a_1; a_2; a_3; z)$ is the Gaussian hypergeometric function with $F_1 (a_1; a_2; a_3; z)$. 

5
And as

\[
\frac{\zeta_2}{\zeta_2 - 1} = -\frac{\Lambda_y}{\varphi(\beta, b)} (1 - \hat{\rho}(\beta, b))^2 = \frac{\Lambda_y}{\varphi(\beta, b)} (\hat{\rho}(\beta, b) - 1)^2
\]

Hence

\[
I_2 = \sqrt{\frac{\Lambda_y}{\varphi(\beta, b)}} (\hat{\rho}(\beta, b) - 1) \times 2 F_1 \left( \frac{N(T - 1)}{2}; \frac{1}{2}; \frac{3}{2}; \frac{\Lambda_y}{\varphi(\beta, b)} (\hat{\rho}(\beta, b) - 1)^2 \right)
\]

Then, the denominator of the conditional posterior density of \( \rho \), given \((\beta, b)\), is

\[
A = [\varphi(\beta, b)]^{-\frac{N(T-1)}{2}} \sqrt{\frac{\Lambda_y}{\varphi(\beta, b)}} \left[ I_1 + I_2 \right]
\]

\[
= [\varphi(\beta, b)]^{-\frac{N(T-1)}{2}} \sqrt{\frac{\Lambda_y}{\varphi(\beta, b)}} \left[ \left( \hat{\rho}(\beta, b) - 1 \right) \times 2 F_1 \left( \frac{N(T-1)}{2}; \frac{1}{2}; \frac{3}{2}; \frac{\Lambda_y}{\varphi(\beta, b)} (\hat{\rho}(\beta, b) - 1)^2 \right) \right]
\]

and the conditional posterior density of \( \rho \), given \((\beta, b)\), is

\[
\pi^*(\rho|\beta, b) = \frac{1}{A} \left[ \varphi(\beta, b) + \Lambda_y \{ \rho - \hat{\rho}(\beta, b) \}^2 \right]^{-\frac{N(T-1)}{2}}
\]

\[
= \frac{\varphi(\beta, b)^{-\frac{N(T-1)}{2}}}{(AB)} B \left[ 1 + \frac{\Sigma^{-1}}{N(T-1) - 2} \{ \rho - \hat{\rho}(\beta, b) \}^2 \right]^{-\frac{N(T-1)}{2}}
\]

where\(^2\)

\[
\Sigma^{-1} = \frac{\Lambda_y [N(T-1) - 2]}{\varphi(\beta, b)}
\]

\[
B = \frac{\Gamma \left( \frac{N(T-1)}{2} \right)}{\Gamma \left( \frac{N(T-1)}{2} - 1 \right)} \sqrt{\frac{\Lambda_y}{\varphi(\beta, b)}} = \frac{N(T-1)}{2\pi [N(T-1) - 2]} \sqrt{\frac{\Lambda_y}{\varphi(\beta, b)}}
\]

Then, the expression

\[
B \left[ 1 + \frac{\Sigma^{-1}}{N(T-1) - 2} \{ \rho - \hat{\rho}(\beta, b) \}^2 \right]^{-\frac{N(T-1)}{2}}
\]

is the pdf of a \( t_\nu(\hat{\rho}(\beta, b), \Sigma) \) with \( \Sigma = \frac{\varphi(\beta, b)}{\Lambda_y [N(T-1) - 2]} \) and \( \nu = N(T-1) - 2 \) degrees of freedom. As \( \pi^*(\rho|\beta, b) \) is a linear combination of a \( t_\nu \) distribution (e.g., \( C^{-1} t_\nu \) with

\(^2\) A random variable \( X \in \mathbb{R}^p \) has a multivariate Student distribution with location parameters \( \mu \), shape matrix \( \Sigma \) and \( \nu \) degrees of freedom, \( X \sim t_\nu(\mu, \Sigma) \) if its pdf is given by

\[
\frac{\Gamma \left( \frac{\nu + p}{2} \right)}{\Gamma \left( \frac{\nu}{2} \right) \nu^{p/2} |\Sigma|^{p/2}} \left[ 1 + \frac{1}{\nu} (x - \mu)' \Sigma^{-1} (x - \mu) \right]^{-\frac{\nu + p}{2}}
\]
As the mean (resp. the variance) of a random variables with degrees of freedom, say $S$, we can use the results of Welch (1947) and Fairweather (1972) (see also Witkovský (2004))\(^3\). As the mean (resp. the variance) of a $t_{\nu_S}$ distribution is $\hat{\rho}(\beta, b)$, for $\nu_S > 1$ (resp. $\nu_S > 2$), then the mean of the posterior density of $\rho$ is:

$$\hat{\rho} = D\hat{\rho}(\beta, b)$$

and the variance of the posterior density of $\rho$ is

$$\text{Var}[\hat{\rho}] = D^2 \frac{\nu_S}{\nu_S - 2} \Sigma$$

with

$$\nu_S = 4 + C^2 (N (T - 1) - 6)$$

and

$$D = \sqrt{\frac{\nu_S - 2}{\nu_S} \frac{N (T - 1) - 2}{N (T - 1) - 4}}$$

If $\rho$ is assumed to be $U(-1, 1)$, then we get a three-step approach. For the dynamic specification: $y = \rho y_{-1} + X \beta + W b + u$, we can integrate first with respect to $(\beta, \tau)$ given $b$ and $\rho$, and then, conditional on $\beta$ and $\rho$, we can next integrate with respect to $(b, \tau)$ and last, we can integrate with respect to $\rho$ given $(\beta, b)$.

1. Let $y^* = (y - \rho y_{-1} - W b)$. Derive the conditional ML-II posterior distribution of $\beta$ given the specific effects $b$ and $\rho$ as in the section 2.3.1 of the main text.
2. Let $\tilde{y} = (y - \rho y_{-1} - X \beta)$. Derive the conditional ML-II posterior distribution of $b$ given the coefficients $\beta$ and $\rho$ as in the section 2.3.2 of the main text.
3. Let $y^2 = (y - X \beta - W b)$. Derive the conditional ML-II posterior distribution of $\rho$ given the coefficients $\beta$ and $b$ as in the previous section A.1.

As the mean and variance of $\rho$ are exactly defined, we do not need to introduce an $\varepsilon$-contamination class of prior distributions for $\rho$ at the second stage of the hierarchy. This was initially our first goal. Unfortunately, the results obtained on a Monte Carlo simulation study (see section A.2) provide

\(^3\)If $S = \sum_{j=1}^{k} \lambda_j t_{\nu_j}$ is a weighted linear combination of independent Student’s $t$ random variables with $\nu_j$ ($j = 1, \ldots, k$) degrees of freedom, then $S$ is approximated by the distribution of the $D$ multiple of single Student’s $t$ random variables with $\nu_S$ degrees of freedom, say $D t_{\nu_S}$, where $\nu_S$ and $D > 0$ are to be determined by equating the second and the fourth moments of $D t_{\nu_S}$ to those of $S$. In particular, if $\nu_j > 4$, \(\forall j\), then:

$$\nu_S = 4 + \frac{1}{\sum_{j=1}^{k} \frac{\nu_j}{\nu_j^2}}, \quad D = \sqrt{\frac{\nu_S - 2}{\nu_S} \frac{1}{\sum_{j=1}^{k} \frac{\nu_j^2 - 2}{\nu_j}}}$$

In our case, $k = 1$, $\lambda_1 = \left(\frac{1}{\nu_1}\right)$ with $C = A.B.\varphi(\beta, b)$, and $\nu_1 = N(T - 1) - 2$. So

$$\nu_S = 4 + C^2 (\nu_1 - 4) = 4 + C^2 (N (T - 1) - 6)$$

and

$$D = \sqrt{\frac{\nu_S - 2}{\nu_S} \frac{\nu_1}{\nu_1 - 2}} = \sqrt{\frac{\nu_S - 2}{\nu_S} \frac{N (T - 1) - 2}{N (T - 1) - 4}}$$
biased estimates of $\rho$, $\beta$ and residual variances. That is why we assume a Zellner $g$-prior, for the $\theta(=[\rho, \beta']')$ vector encompassing the coefficient of the lagged dependent variable $y_{i,t-1}$ and those of the explanatory variables $X'_{it}$. Thus, we do not impose stationarity constraints like many authors and we respect the philosophy of $\varepsilon$-contamination class using data-driven priors.

A.2. Some Monte Carlo results

We run a Monte Carlo simulation study for the dynamic random effects world comparing different robust Bayesian estimators. As previously in the main text, we run the two stage approach with individual block resampling bootstrap assuming a Zellner $g$-prior, for the $\theta(=[\rho, \beta']')$ vector encompassing the coefficient of the lagged dependent variable $y_{i,t-1}$ and those of the explanatory variables $X'_{it}$. We introduce a two stage three step approach when $\rho \sim U(-1, 1)$. When the initial value of $\rho$ is drawn for a uniform distribution $U(-1, 1)$, results are strongly biased as shown on Table A.1. Even if we initialize $\rho$ with its OLS estimator on the pooled model, the results, if they improve, are further biased.
Table A.1: Dynamic Random Effects World

$N = 100$, $T = 10$, $\varepsilon = 0.5$, Replications=1,000

<table>
<thead>
<tr>
<th></th>
<th>$\rho$</th>
<th>$\beta_{11}$</th>
<th>$\beta_{12}$</th>
<th>$\beta_{2}$</th>
<th>$\sigma^2_u$</th>
<th>$\sigma^2_\mu$</th>
<th>$\lambda_\theta$</th>
<th>$\lambda_\mu$</th>
<th>Computation Time (secs.)</th>
</tr>
</thead>
<tbody>
<tr>
<td>true</td>
<td>0.75</td>
<td>1.0252</td>
<td>0.9952</td>
<td>1.0207</td>
<td>0.9976</td>
<td>4.0433</td>
<td>&lt; $10^{-4}$</td>
<td>0.4943</td>
<td>521.89</td>
</tr>
<tr>
<td>2S boot</td>
<td>coef</td>
<td>0.7522</td>
<td>1.0252</td>
<td>0.9952</td>
<td>1.0207</td>
<td>0.9976</td>
<td>4.0433</td>
<td>&lt; $10^{-4}$</td>
<td>0.4943</td>
</tr>
<tr>
<td></td>
<td>se</td>
<td>0.0060</td>
<td>0.0249</td>
<td>0.0253</td>
<td>0.0252</td>
<td>0.0481</td>
<td>0.2025</td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>rmse</td>
<td>0.0055</td>
<td>0.0351</td>
<td>0.0248</td>
<td>0.0324</td>
<td>0.0482</td>
<td>0.2070</td>
<td></td>
<td></td>
</tr>
<tr>
<td>2S-3S boot</td>
<td>coef</td>
<td>0.2358</td>
<td>1.2131</td>
<td>1.1870</td>
<td>1.2510</td>
<td>5.4548</td>
<td>167.5906</td>
<td>0.0527</td>
<td>0.3821</td>
</tr>
<tr>
<td>$U(-1, 1)$</td>
<td>se</td>
<td>0.3397</td>
<td>0.1923</td>
<td>0.1970</td>
<td>0.2063</td>
<td>0.5202</td>
<td>0.2412</td>
<td>0.2159</td>
<td>0.2749</td>
</tr>
<tr>
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<td>rmse</td>
<td>0.5202</td>
<td>0.2412</td>
<td>0.2159</td>
<td>0.2749</td>
<td>0.5202</td>
<td>0.2412</td>
<td>0.2159</td>
<td>0.2749</td>
</tr>
<tr>
<td>2S-3S boot</td>
<td>coef</td>
<td>0.7379</td>
<td>1.1174</td>
<td>1.0764</td>
<td>1.1148</td>
<td>1.0400</td>
<td>4.6650</td>
<td>0.0465</td>
<td>0.4920</td>
</tr>
<tr>
<td>$U(-1, 1)$ init OLS</td>
<td>se</td>
<td>0.0065</td>
<td>0.0275</td>
<td>0.0273</td>
<td>0.0282</td>
<td>0.0131</td>
<td>0.1200</td>
<td>0.0801</td>
<td>0.1175</td>
</tr>
<tr>
<td></td>
<td>rmse</td>
<td>0.0131</td>
<td>0.1200</td>
<td>0.0801</td>
<td>0.1175</td>
<td>0.0131</td>
<td>0.1200</td>
<td>0.0801</td>
<td>0.1175</td>
</tr>
</tbody>
</table>

2S bootstrap: two stage with individual block resampling bootstrap.

2S-3S boot $U(0, 1)$: two stage - three steps with individual block resampling bootstrap and with $\rho \sim U(-1, 1)$

where $\rho$ init is drawn from $U(-1, 1)$.

2S-3S boot $U(0, 1)$ init OLS: two stage - three steps with individual block resampling bootstrap and with $\rho \sim U(-1, 1)$

where $\rho$ init is the OLS estimator of $\rho$. 


B. Some extra Monte Carlo simulation results

1. Dynamic random effects world with $\rho = 0.98$.
2. Chamberlain-type Fixed Effects World with $\rho = 0.75$.
3. Sensitivity to $\varepsilon$-contamination values
4. Departure from normality
Table B.2: Dynamic Random Effects World

\(N = 100, T = 10, \varepsilon = 0.5, \text{Replications}=1,000\)

<table>
<thead>
<tr>
<th>(\rho)</th>
<th>(\beta_{11})</th>
<th>(\beta_{12})</th>
<th>(\beta_{2})</th>
<th>(\sigma_u^2)</th>
<th>(\sigma_{\mu}^2)</th>
<th>(\lambda_\theta)</th>
<th>(\lambda_\mu)</th>
<th>Computation Time (secs.)</th>
</tr>
</thead>
<tbody>
<tr>
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<td>1</td>
<td>1</td>
<td>1</td>
<td>1</td>
<td>1</td>
<td>1</td>
<td>1</td>
</tr>
<tr>
<td>2S boot</td>
<td>coef</td>
<td>0.9822</td>
<td>1.0151</td>
<td>0.9843</td>
<td>1.0087</td>
<td>0.9971</td>
<td>3.7994</td>
<td>(&lt;10^{-4})</td>
</tr>
<tr>
<td></td>
<td>se</td>
<td>0.0009</td>
<td>0.0242</td>
<td>0.0247</td>
<td>0.0250</td>
<td>0.0485</td>
<td>0.2003</td>
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</tr>
<tr>
<td></td>
<td>rmse</td>
<td>0.0022</td>
<td>0.0339</td>
<td>0.0334</td>
<td>0.0306</td>
<td>0.0485</td>
<td>0.2833</td>
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<tr>
<td></td>
<td>hpdi_lower</td>
<td>0.9801</td>
<td>0.9482</td>
<td>0.9185</td>
<td>0.9365</td>
<td>0.9016</td>
<td>3.4361</td>
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<td>hpdi_upper</td>
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<td>1.0832</td>
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<td>1.0766</td>
<td>1.0908</td>
<td>4.1940</td>
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<tr>
<td>GMM</td>
<td>coef</td>
<td>0.9831</td>
<td>0.9300</td>
<td>0.8502</td>
<td>0.9279</td>
<td>0.9016</td>
<td>4.0194</td>
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</tr>
<tr>
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<td>se</td>
<td>0.0009</td>
<td>0.0282</td>
<td>0.0273</td>
<td>0.0290</td>
<td>0.0561</td>
<td>0.4290</td>
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</tr>
<tr>
<td></td>
<td>rmse</td>
<td>0.0036</td>
<td>0.0857</td>
<td>0.1568</td>
<td>0.0879</td>
<td>0.1132</td>
<td>0.4292</td>
<td></td>
</tr>
<tr>
<td>QMLE</td>
<td>coef</td>
<td>0.9794</td>
<td>1.0110</td>
<td>0.9771</td>
<td>1.0083</td>
<td>0.9942</td>
<td>4.1953</td>
<td></td>
</tr>
<tr>
<td></td>
<td>se</td>
<td>0.0007</td>
<td>0.0256</td>
<td>0.0252</td>
<td>0.0255</td>
<td>0.0508</td>
<td>0.2606</td>
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<tr>
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<td>0.0279</td>
<td>0.0279</td>
<td>0.0279</td>
<td>0.0511</td>
<td>0.3255</td>
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</table>

2S boot: two stage with individual block resampling bootstrap.
GMM: Arellano-Bond GMM estimation.
QMLE: quasi-maximum likelihood estimation.
hpdi\_lower (hpdi\_upper): lower and upper bounds of the 95% Highest Posterior Density Interval (HPDI).
### Table B.3: Dynamic Chamberlain-type Fixed Effects World

$N = 100, T = 10, \varepsilon = 0.5, \text{Replications}=1,000$

<table>
<thead>
<tr>
<th></th>
<th>$\rho$</th>
<th>$\beta_{11}$</th>
<th>$\beta_{12}$</th>
<th>$\beta_2$</th>
<th>$\sigma_u^2$</th>
<th>$\sigma_p^2$</th>
<th>$\lambda_0$</th>
<th>$\lambda_\mu$</th>
<th>Computation Time (secs.)</th>
</tr>
</thead>
<tbody>
<tr>
<td>true</td>
<td>0.75</td>
<td>1</td>
<td>1</td>
<td>1</td>
<td>1</td>
<td>1</td>
<td>1</td>
<td></td>
<td>162.5341</td>
</tr>
<tr>
<td>2S boot coef</td>
<td>0.7414</td>
<td>1.0134</td>
<td>1.0132</td>
<td>0.9959</td>
<td>0.9952</td>
<td>173.8813</td>
<td>&lt; $10^{-4}$</td>
<td>0.4959</td>
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<tr>
<td>se</td>
<td>0.0061</td>
<td>0.0234</td>
<td>0.0236</td>
<td>0.0231</td>
<td>0.0665</td>
<td>36.1240</td>
<td></td>
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</tr>
<tr>
<td>rmse</td>
<td>0.0105</td>
<td>0.0277</td>
<td>0.0274</td>
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<td>0.0666</td>
<td>37.8470</td>
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<table>
<thead>
<tr>
<th></th>
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<th>$\pi_5$</th>
<th>$\pi_6$</th>
<th>$\pi_7$</th>
<th>$\pi_8$</th>
<th>$\pi_9$</th>
<th>$\pi_{10}$</th>
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</thead>
<tbody>
<tr>
<td>true</td>
<td>0.1678</td>
<td>0.2097</td>
<td>0.2621</td>
<td>0.3277</td>
<td>0.4096</td>
<td>0.5120</td>
<td>0.6400</td>
<td>0.8000</td>
<td>1.0000</td>
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<tr>
<td>2S boot coef</td>
<td>0.1865</td>
<td>0.2267</td>
<td>0.2812</td>
<td>0.3465</td>
<td>0.4284</td>
<td>0.5327</td>
<td>0.6622</td>
<td>0.8224</td>
<td>1.0259</td>
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<tr>
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<td>0.1068</td>
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<td>0.1081</td>
<td>0.1072</td>
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<td>0.1089</td>
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<td>rmse</td>
<td>0.0797</td>
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<th>$\beta_{12}$</th>
<th>$\beta_2$</th>
<th>$\sigma_u^2$</th>
<th>$\sigma_p^2$</th>
<th>$\lambda_0$</th>
<th>$\lambda_\mu$</th>
<th>Computation Time (secs.)</th>
</tr>
</thead>
<tbody>
<tr>
<td>QMLE coef</td>
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<td>1199.07</td>
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<tr>
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<table>
<thead>
<tr>
<th></th>
<th>$\pi_2$</th>
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<th>$\pi_4$</th>
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<th>$\pi_9$</th>
<th>$\pi_{10}$</th>
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</thead>
<tbody>
<tr>
<td>true</td>
<td>0.1678</td>
<td>0.2097</td>
<td>0.2621</td>
<td>0.3277</td>
<td>0.4096</td>
<td>0.5120</td>
<td>0.6400</td>
<td>0.8000</td>
<td>1.0000</td>
</tr>
<tr>
<td>QMLE coef</td>
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<td>0.2616</td>
<td>0.3274</td>
<td>0.4083</td>
<td>0.5090</td>
<td>0.6388</td>
<td>0.7958</td>
<td>0.9961</td>
</tr>
<tr>
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<td>0.0807</td>
<td>0.0858</td>
<td>0.0888</td>
<td>0.0971</td>
</tr>
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<td>0.0782</td>
<td>0.0759</td>
<td>0.0810</td>
<td>0.0807</td>
<td>0.0857</td>
<td>0.0888</td>
<td>0.0971</td>
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</tbody>
</table>

2S boot: two stage with individual block resampling bootstrap.
QMLE: quasi-maximum likelihood estimation.
Table B.4: Dynamic Chamberlain-type Fixed Effects World

\(N = 200, T = 30, \varepsilon = 0.5\), Replications=500(*)

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<th></th>
<th>(\rho)</th>
<th>(\beta_{11})</th>
<th>(\beta_{12})</th>
<th>(\beta_{2})</th>
<th>(\sigma_{u}^2)</th>
<th>(\sigma_{\mu}^2)</th>
<th>(\lambda_0)</th>
<th>(\lambda_\mu)</th>
<th>Computation Time (secs.)</th>
</tr>
</thead>
<tbody>
<tr>
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<td>1</td>
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<td>1</td>
<td>1</td>
<td>1</td>
<td></td>
<td>204.3142</td>
</tr>
<tr>
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<td>1.0059</td>
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<td>1.0005</td>
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<td>&lt; 10^{-4}</td>
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</tr>
<tr>
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<td>0.0087</td>
<td>0.0091</td>
<td>0.0267</td>
<td>28.4166</td>
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<td></td>
<td></td>
</tr>
<tr>
<td>rmse</td>
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<td>0.0110</td>
<td>0.0107</td>
<td>0.0094</td>
<td>0.0266</td>
<td>28.7231</td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

\[
\begin{align*}
\pi_2 & = 0.0019 \\
\pi_3 & = 0.0024 \\
\pi_4 & = 0.0030 \\
\pi_5 & = 0.0038 \\
\pi_6 & = 0.0047 \\
\pi_7 & = 0.0059 \\
\pi_8 & = 0.0074 \\
\pi_9 & = 0.0092 \\
\pi_{10} & = 0.0115
\end{align*}
\]

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<thead>
<tr>
<th></th>
<th>(\pi_{11})</th>
<th>(\pi_{12})</th>
<th>(\pi_{13})</th>
<th>(\pi_{14})</th>
<th>(\pi_{15})</th>
<th>(\pi_{16})</th>
<th>(\pi_{17})</th>
<th>(\pi_{18})</th>
<th>(\pi_{19})</th>
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<td>0.0225</td>
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\pi_{24} & = 0.0426 \\
\pi_{25} & = 0.0599 \\
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\pi_{27} & = 0.0876 \\
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\end{align*}
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(*) When \(T = 30\), we restrict the exercise to only 500 replications, not because of our estimator under R but because of the size of the simulated database and its reading under Stata. Indeed, for 1,000 replications and \(T = 30\), one must read, under Stata, \((1 + 4 + 30) \times 1,000 = 35,000\) variables of size \(NT, 1\)! Even with a 64-bit computer and Stata (MP and S), only 32,767 variables can be read. On the other hand, with our code under R, there is no limitation of that order.
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### Table B.5: Dynamic Random Effects World, robustness to $\varepsilon$-contamination

$N = 100, T = 10, \text{replications}=1,000$

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2S boot: two stage with individual block resampling bootstrap.
GMM: Arellano-Bond GMM estimation.
QMLE: quasi-maximum likelihood estimation.
Table B.6: Dynamic heterogeneous panel data model with common correlated effects, robustness to ε-contamination

\( N = 100, T = 30, \text{replications}=1,000 \)

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<th>( \lambda_\theta )</th>
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<td>0.0248</td>
<td>0.0882</td>
<td>0.0926</td>
<td>0.0664</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>( \varepsilon = 0.9 )</td>
<td>2S boot coef</td>
<td>0.7347</td>
<td>0.8071</td>
<td>1.0759</td>
<td>1.0536</td>
<td>0.0010</td>
<td>0.0738</td>
</tr>
<tr>
<td>se</td>
<td>0.0194</td>
<td>0.0671</td>
<td>0.0531</td>
<td>0.0392</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>mse</td>
<td>0.0248</td>
<td>0.0882</td>
<td>0.0926</td>
<td>0.0664</td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

2S boot: two stage with individual block resampling bootstrap.
CCEMG: Common Correlated Effects Mean Group estimator.
Mean coefficients: \( \bar{\rho} = (1/N) \sum_{i=1}^{N} \rho_i \) and \( \bar{\beta}_1 = (1/N) \sum_{i=1}^{N} \beta_{1i} \).
Table B.7: Dynamic Random Effects World, robustness to $\varepsilon$-contamination, departure from normality: the skewed $t$-distribution

$N = 100$, $T = 10$, $\varepsilon = 0.5$, replications=1,000

<table>
<thead>
<tr>
<th>$\rho$</th>
<th>$\beta_{11}$</th>
<th>$\beta_{12}$</th>
<th>$\beta_2$</th>
<th>$\sigma_u^2$</th>
<th>$\sigma_\mu^2$</th>
<th>$\lambda_\theta$</th>
<th>$\lambda_\mu$</th>
<th>Computation Time (secs.)</th>
</tr>
</thead>
<tbody>
<tr>
<td>true</td>
<td>0.75</td>
<td>1</td>
<td>1</td>
<td>1</td>
<td>7.0052</td>
<td>4</td>
<td></td>
<td>149.15</td>
</tr>
<tr>
<td>GMM coef</td>
<td>0.7588</td>
<td>0.9871</td>
<td>0.8901</td>
<td>0.9824</td>
<td>6.0976</td>
<td>5.9883</td>
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<td></td>
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<tr>
<td>se</td>
<td>0.0104</td>
<td>0.0449</td>
<td>0.0414</td>
<td>0.0456</td>
<td>4.5903</td>
<td>1.1453</td>
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</tr>
<tr>
<td>rmse</td>
<td>0.0196</td>
<td>0.0683</td>
<td>0.1262</td>
<td>0.0698</td>
<td>4.6769</td>
<td>2.2943</td>
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<td></td>
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<tr>
<td>2S boot coef</td>
<td>0.7141</td>
<td>1.0711</td>
<td>1.0382</td>
<td>1.0714</td>
<td>6.7171</td>
<td>5.8258</td>
<td>0.0004</td>
<td>0.4984</td>
</tr>
<tr>
<td>se</td>
<td>0.0169</td>
<td>0.0625</td>
<td>0.0625</td>
<td>0.0651</td>
<td>4.0849</td>
<td>1.8050</td>
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<td></td>
</tr>
<tr>
<td>rmse</td>
<td>0.0402</td>
<td>0.0956</td>
<td>0.0756</td>
<td>0.0985</td>
<td>4.0931</td>
<td>2.5668</td>
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</tr>
<tr>
<td>QMLE coef</td>
<td>0.7471</td>
<td>1.0215</td>
<td>0.9414</td>
<td>1.0172</td>
<td>6.8577</td>
<td>4.0629</td>
<td>0.0004</td>
<td>0.4984</td>
</tr>
<tr>
<td>se</td>
<td>0.0158</td>
<td>0.0637</td>
<td>0.0625</td>
<td>0.0631</td>
<td>4.8017</td>
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<tr>
<td>rmse</td>
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<td>0.0673</td>
<td>0.0673</td>
<td>4.8016</td>
<td>0.7317</td>
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</table>

2S boot: two stage with individual block resampling bootstrap.
GMM: Arellano-Bond GMM estimation.
QMLE: quasi-maximum likelihood estimation.

Table B.8: Dynamic heterogeneous panel data model with common correlated effects, departure from normality: the skewed $t$-distribution

$N = 100$, $T = 30$, $\varepsilon = 0.5$, replications=1,000

<table>
<thead>
<tr>
<th>$\bar{\rho}$</th>
<th>$\bar{\beta}_1$</th>
<th>$\bar{\beta}_2$</th>
<th>$\sigma_u^2$</th>
<th>$\lambda_\theta$</th>
<th>$\lambda_\mu$</th>
<th>Computation Time (secs.)</th>
</tr>
</thead>
<tbody>
<tr>
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<td>0.7501</td>
<td>0.7498</td>
<td>1</td>
<td>6.9824</td>
<td></td>
<td></td>
</tr>
<tr>
<td>CCEMG coef</td>
<td>0.6300</td>
<td>0.8064</td>
<td>1.2112</td>
<td>7.3744</td>
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</tr>
<tr>
<td>se</td>
<td>0.0319</td>
<td>0.1364</td>
<td>0.1119</td>
<td>1.6597</td>
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<td></td>
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<tr>
<td>rmse</td>
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<td>0.1477</td>
<td>0.2390</td>
<td>6.5867</td>
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<tr>
<td>2S boot coef</td>
<td>0.6930</td>
<td>0.8788</td>
<td>1.2188</td>
<td>7.0220</td>
<td>0.0110</td>
<td>0.4202</td>
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<tr>
<td>se</td>
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<td>0.1292</td>
<td>0.1018</td>
<td>2.5438</td>
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<td>0.1825</td>
<td>0.2413</td>
<td>2.5428</td>
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</tr>
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</table>

2S boot: two stage with individual block resampling bootstrap.
CCEMG: Common Correlated Effects Mean Group estimator.
mean coefficients: $\bar{\rho} = (1/N) \sum_{i=1}^N \rho_i$ and $\bar{\beta}_1 = (1/N) \sum_{i=1}^N \beta_{1i}$. 
References


Welch, B.L., 1947. The generalization of Student’s problem when several different population variances are involved. Biometrika 34, 28–35.