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Large Bayesian VARMAs*

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Abstract: Vector Autoregressive Moving Average (VARMA) models have many theoretical properties which should make them popular among empirical macroeconomists. However, they are rarely used in practice due to over-parameterization concerns, difficulties in ensuring identification and computational challenges. With the growing interest in multivariate time series models of high dimension, these problems with VARMAs become even more acute, accounting for the dominance of VARs in this field. In this paper, we develop a Bayesian approach for inference in VARMAs which surmounts these problems. It jointly ensures identification and parsimony in the context of an efficient Markov chain Monte Carlo (MCMC) algorithm. We use this approach in a macroeconomic application involving up to twelve dependent variables. We find our algorithm to work successfully and provide insights beyond those provided by VARs.

Keywords: VARMA identification, Markov Chain Monte Carlo, Bayesian, stochastic search variable selection

JEL Classification: C11, C32, E37

1   Introduction

Vector autoregressions (VARs) have been extremely popular in empirical macroeconomics and other fields for several decades (e.g. beginning with early work such as Sims, 1980, Doan, Litterman and Sims, 1984 and Litterman, 1986 with recent examples being Korobilis, 2013 and Koop, 2014). Until recently, most of these VARs have involved only a few (e.g. two to seven) dependent variables. However, VARs involving tens or even hundreds of variables are increasingly popular (see, e.g., Banbura, Giannone and Reichlin, 2010, Carriero, Clark and Marcellino, 2011, Carriero, Kapetanios and Marcellino, 2009, Giannone, Lenza, Momferatou and Onorante, 2010 and Koop, 2013, and Gefang, 2014). Vector autoregressive moving average models (VARMAs) have enjoyed less popularity with empirical researchers despite the fact that theoretical macroeconomic models such as dynamic stochastic general equilibrium models (DSGEs) lead to VMA representations which may not be well approximated by VARs, especially parsimonious VARs with short lag lengths. Papers such as Cooley and Dwyer (1998) point out the limitations of the structural VAR (SVAR) framework and suggest VARMA models as often being more appropriate. For instance, Cooley and Dwyer (1998) conclude "While VARMA models involve additional estimation and identification issues, these complications do not justify systematically ignoring these moving average components, as in the SVAR approach." There is, thus, a strong justification for the empirical macreconomist's toolkit to include VARMAs. Papers such as Poskitt and Yao (2012) document the errors which arise when approximating a VARMA with a finite order VAR and show them to be potentially substantial.[1]

VARs are commonly used for forecasting. But, for the forecaster, too, there are strong reasons to be interested in VARMAs. The univariate literature contains numerous examples in finance and macroeconomics where adding MA components to AR models improves forecasting (e.g. Chan, 2013). But even with multivariate macroeconomic forecasting some papers (e.g. Athanasopoulos and Vahid, 2008) have found that VARMAs forecast better than VARs. Theoretical econometric papers such as Lutkepohl and Poskitt (1996) point out further advantages of VARMAs over VARs.

Despite these advantages of VARMA models, they are rarely used in practice. There are three main reasons for this. First, there are difficult identification problems to be overcome. Second, VARMAs are parameter rich models which can be over-parameterized (an especially important concern in light of the growing interest in large dimensional models as is evinced in the large VAR literature). And, largely due to the first two problems, they can be difficult to estimate. This paper develops methods for estimating VARMAs which address all these concerns.

The paper is organized in the following sections. Section 2 describes the econometric theory of VARMAs paying particular attention to different parameterizations of the VARMA including the expanded form (which is used in the main part of our MCMC algorithm) and the echelon form (which is used in our treatment of identification). Section 3 describes our approach which uses Bayesian methods and a hierarchical prior to jointly

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[1]Poskitt and Yao (2012) also show that, asymptotically, the error involved in this approximation vanishes far more slowly than estimation error.

select identification restrictions and ensure shrinkage in the resulting model. An MCMC algorithm which implements our approach is developed. Section 4 investigates how well our approach works in practice through a substantive macroeconomic application using VARMAs containing up to 12 variables. We find that our methods are computationally feasible and lead to inference on parameters and impulse responses that are more reasonable and estimated more precisely than alternative approaches, especially in the larger VARMAs of interest in modern macroeconomics. An online appendix, available at http://personal.strath.ac.uk/gary.koop/research.htm, contains additional technical details and empirical results as well as an empirical exercise using artificially generated data.

2 The Econometrics of VARMAs

2.1 The Semi-structural VARMA

Consider the n dimensional multivariate time series $\mathbf{y}_t, t = -\infty, \ldots, \infty$ and begin with the semi-structural form of the VARMA(p, q):

$$\mathbf{B}_0 \mathbf{y}_t = \sum_{j=1}^{p} \mathbf{B}_j \mathbf{y}_{t-j} + \sum_{j=1}^{q} \boldsymbol{\Theta}_j \boldsymbol{\epsilon}_{t-j} + \boldsymbol{\Theta}_0 \boldsymbol{\epsilon}_t, \qquad \boldsymbol{\epsilon}_t \sim \mathcal{N}(\mathbf{0}, \boldsymbol{\Sigma}) \tag{1}$$

or, in terms of matrix polynomial lag operators,

$$\mathbf{B}(L)\mathbf{y}_t = \boldsymbol{\Theta}(L)\boldsymbol{\epsilon}_t,$$

and assume stationarity and invertibility.[2] For future reference, denote the elements of the VAR and VMA parts of the model as $\mathbf{B}(L) = [B_{ik}(L)]$ and $\boldsymbol{\Theta}(L) = [\Theta_{ik}(L)]$ for $i, k = 1, .., n$.

The theoretical motivation for the VARMA arises from the Wold decomposition:

$$\mathbf{y}_t = \mathbf{K}(L)\boldsymbol{\epsilon}_t, \tag{2}$$

where $\mathbf{K}(L)$ is generally an infinite degree polynomial operator. Specifically, it can be shown that any such rational transfer function $\mathbf{K}(L)$ corresponds to the existence of two finite degree operators $\mathbf{B}(L)$ and $\boldsymbol{\Theta}(L)$ such that

$$\mathbf{B}(L)\mathbf{K}(L) = \boldsymbol{\Theta}(L).$$

Thus, the VARMA(p, q) is an exact finite-order representation of any multivariate system that can be characterized by a rational transfer function. When $\mathbf{K}(L)$ is not rational, the VARMA(p, q) can provide an arbitrarily close approximation. Moreover, an important advantage of the VARMA class is that, unlike VARs or pure VMAs, it is closed under a variety of transformations of $\mathbf{y}_t$, including linear operations and subsets.

The practical problem in having both AR terms with MA terms, however, is that an alternative VARMA with coefficients $\mathbf{B}^\dagger(L) = \mathbf{C}(L)\mathbf{B}(L)$ and $\boldsymbol{\Theta}^\dagger(L) = \mathbf{C}(L)\boldsymbol{\Theta}(L)$

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[2]In principle, our algorithm would work with non-stationary data, although priors may have to be adjusted relative to the choices we make.

will lead to the same Wold representation. The VARMA(p, q) representation, therefore, is in general not unique. However, there are two reasons why a unique representation is desirable in practice: parsimony and identification. The first reason concerns both frequentist and Bayesian approaches. If $\mathbf{B}(L)$ and $\mathbf{\Theta}(L)$ contain redundancies, then the resulting model may lead to poor forecast performance and imprecise impulse response functions. For researchers working with larger VARMAs such over-parameterization concerns can become severe. For instance, in our empirical work, we use as an estimating model the 12-variate VARMA with four lags (and an intercept in each equation). Even if the conventional restriction that $\mathbf{B}_0 = \mathbf{\Theta}_0 = \mathrm{I}$ is imposed,[3] there are still 1,242 parameters (including error covariances) to estimate. With macroeconomic data sets containing a few hundred observations, it will be very hard to obtain precise inference for all these parameters in the absence of an econometric method which ensures parsimony or shrinkage.

The second reason (lack of identification) may be less important for the Bayesian who is only interested in forecasting or in identified functions of the parameters such as impulse responses. That is, given a proper prior a well-defined posterior will exist even in a non-identified VARMA. However, the role of the prior becomes important in such cases and carelessly constructed priors can lead to deficient inference for the Bayesian. And, for the Bayesian interested in issues such as Granger causality and weak exogeneity or working with a VARMA as an approximation to a DSGE model, it is typically useful to work with an identified model. For frequentists, however, a lack of identification is a more substantive problem, precluding estimation.

How does one obtain a unique VARMA representation? There are generally two major steps:

The first step is to eliminate common roots in $\mathbf{B}(L), \mathbf{\Theta}(L)$ such that only $\mathbf{C}(L)$ with a constant determinant is possible. In this case, the operators $\mathbf{B}(L), \mathbf{\Theta}(L)$ are said to be left coprime and $\mathbf{C}(L)$ unimodular. For the univariate case, it is sufficient to achieve uniqueness and corresponds in practical terms to specifying minimal orders p, q. For a multivariate process, however, this is not enough and a second step is required. That is, even if $\mathbf{B}_0 = \mathbf{\Theta}_0 = \mathrm{I}$ is imposed, there may still exist $\mathbf{C}(L) \neq \mathrm{I}$ that preserves this restriction for an alternative set of left coprime operators $\mathbf{B}^\dagger(L), \mathbf{\Theta}^\dagger(L)$. A common example is

$$\mathbf{C}(L) = \begin{pmatrix} 1 & c(L) \\ 0 & 1 \end{pmatrix}.$$

Clearly, $\det \mathbf{C}(L) = 1$ and for any $\mathbf{B}(L), \mathbf{\Theta}(L)$, the transformations $\mathbf{B}^\dagger(L) = \mathbf{C}(L)\mathbf{B}(L)$ and $\mathbf{\Theta}^\dagger(L) = \mathbf{C}(L)\mathbf{\Theta}(L)$ lead to $\mathbf{B}_0^\dagger = \mathbf{\Theta}_0^\dagger = \mathrm{I}$.

This implies that the elements of $\mathbf{B}(L), \mathbf{\Theta}(L)$ are not identified. One approach to achieving identification relies on the assumption that the matrix $[\mathbf{B}_p : \mathbf{\Theta}_q]$ has full row rank, and indeed, when this holds then $\mathbf{B}_0 = \mathbf{\Theta}_0 = \mathrm{I}$ induces a unique representation (e.g., Hannan, 1976). In practice, one could try to explicitly enforce $[\mathbf{B}_p : \mathbf{\Theta}_q]$ to have full row rank, but that may not be desirable in many applications. The full row rank condition will likely not be satisfied by most data generating processes (DGPs) in practice

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[3]Most estimation procedures for the VARMA impose $\mathbf{B}_0 = \mathbf{\Theta}_0 = \mathrm{I}$ and, for this reason, we refer to this as the conventional restriction. The echelon form used in this paper does not impose this restriction.

(Lütkepohl and Poskitt, 1996). Therefore, forcing it in an estimation routine would likely result in mis-specification and an alternative second step would be required to achieve uniqueness when $[\mathbf{B}_p : \mathbf{\Theta}_q]$ is rank deficient.

The more general approach that we follow involves imposing exclusion restrictions on elements of $\mathbf{B}(L), \mathbf{\Theta}(L)$ such that only $\mathbf{C}(L) = \mathbf{I}$ is possible. It turns out that when such zero restrictions are applied according to a specific set of rules, it is possible to achieve a unique VARMA representation corresponding to a particular rational $\mathbf{K}(L)$. This leads to the echelon form which we will use as a basis for our approach to identification.

We stress that in this paper we are focussing only on statistical identification in the VARMA where a lack of statistical identification is associated with multiple values of the parameters yielding the same value for the likelihood function. We are not contributing to the literature on identification issues relating to the underlying economic structure. To put this another way, in the VAR literature, it is common to use structural VAR models which involve identification restrictions needed to provide a unique mapping from the reduced form VAR parameters to the parameters of a structural economic model. Papers such as Rubio-Ramirez, Waggoner and Zha (2009) establish conditions for identification in structural VARs. In this paper, we are not attempting to do something similar for structural VARMAs.

2.2 The Echelon Form for the VARMA

There are several alternatives to the echelon form when it comes to imposing identification in the VARMA (see, for instance, Chapter 12 of Lutkepohl, 2005). However, as we shall see, the echelon form is both flexible and parsimonious. It is flexible in the sense that any VARMA has an echelon form representation. This contrasts with some otherwise attractive representations, such as that of Zadrozny (2014).[4] An alternative approach to canonical specification and estimation of VARMAs is the scalar component model (SCM), as introduced by Tiao and Tsay (1989), Tsay (1989, 1991) and extended by Athanasopoulos and Vahid (2008). In fact, it has been argued (e.g. Tsay, 1989) that this approach could uncover addition zero restrictions in a VARMA, without losing generality, if for example the true VARMA has MA lag orders that differ from AR lag orders. However, Athanasopolous et al (2012) have shown that the underlying structure of SCM is equivalent to that of an echelon form, and the additional restrictions it uncovers are those that are supported by the data, rather than being necessary for identification. We note that in our Bayesian approach to estimating VARMAs in echelon form, shrinkage priors on VARMA coefficients will also uncover any such additional restrictions in a data-based fashion, such that the resulting specification need not have equal AR and MA lag orders. Unlike working with the echelon form, on the other hand, the method of SCM is difficult to automate as it requires substantial user intervention at various steps, and therefore, it cannot be easily adapted in a Bayesian setting.

The echelon form is parsimonious in that it typically leads to identified VARMAs with fewer parameters than other flexible identification schemes. For instance, Lutkepohl

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[4]This approach assumes controllability which is another kind of potentially restrictive rank restriction on the VARMA coefficients.

(2005) discusses two identification schemes, the echelon form and the final equations form and argues (page 455): "The reader may wonder why we consider the complicated looking echelon representation although the final equations form serves the same purpose. The reason is that the echelon form is usually preferable in practice because it often implies fewer free parameters than the equivalent final equations form."[5] For this reason we focus on the echelon form in this paper.

The echelon form involves a particular set of restrictions on the semi-structural VARMA. The derivation of the echelon form is based on Kronecker index theory which shows that every $\mathbf{K}(L)$ in (2) is associated with a unique set of indices $\boldsymbol{\kappa} = (\kappa_1, \ldots, \kappa_n)$, which can be directly related to the VARMA operators $\mathbf{B}(L), \boldsymbol{\Theta}(L)$. Identification is achieved by imposing restrictions on the VARMA coefficients in (1) according to so-called Kronecker indices $\kappa_1, \ldots, \kappa_n$, with $0 \leq \kappa_i \leq p^*$, where $p^* = \max\{p, q\}$.

To explain further the identifying restrictions in the echelon form note that, without loss of generality, we can denote the VARMA(p, q) as VARMA(p^*, p^*). Then any VARMA(p^*, p^*) can be represented in echelon form by setting $\mathbf{B}_0 = \boldsymbol{\Theta}_0$ to be lower triangular with ones on the diagonal and applying the exclusion restrictions defined by $\boldsymbol{\kappa}$ to $\mathbf{B}_0, \ldots, \mathbf{B}_{p^*}, \boldsymbol{\Theta}_1, \ldots, \boldsymbol{\Theta}_{p^*}$. The latter impose on $[\mathbf{B}(L) : \boldsymbol{\Theta}(L)]$ a maximal degree of each row i equivalent to κ_i plus some additional restrictions specified in the following definition. A VARMA in echelon form is denoted VARMA$_E(\boldsymbol{\kappa})$ and details regarding the foregoing restrictions are discussed in many places. The key theoretical advantage of the echelon form is that, given $\boldsymbol{\kappa}$, it provides a way of constructing a parsimonious VARMA representation for $\mathbf{y}_t$. A by-product of this is that the unrestricted parameters are identified. At the same time, every conceivable VARMA can be represented in echelon form. The formal definition of the echelon form is given, e.g., in Lutkepohl, 2005, page 453 as:

Definition:

The VARMA representation in (1) is in echelon form if the VAR and VMA operators are left coprime and satisfy the following conditions.

The VAR operator is restricted as (for $i, k = 1, \ldots, n$):

$$
\begin{array}{l}
B_{ii}(L) = 1 - \sum_{j=1}^{p_i} B_{j,ii} L^j \text{ for } i = 1, \ldots, n \\
B_{ik}(L) = -\sum_{j=p_i-p_{ik}+1}^{p_i} B_{j,ik} L^j \text{ for } i \neq k
\end{array},
$$

where

$$
p_{ik} = \left\{
\begin{array}{l}
\min(p_i + 1, p_k) \text{ for } i \geq k \\
\min(p_i, p_k) \text{ for } i < k
\end{array}
\right.,
$$

and $\mathbf{B}_0$ is lower triangular with ones on the diagonal. The VMA operator is restricted as (for $i, k = 1, \ldots, n$):

$$
\Theta_{ik}(L) = \sum_{j=0}^{p_i} \Theta_{j,ik} L^j \text{ and } \boldsymbol{\Theta}_0 = \mathbf{B}_0.
$$

The row degrees of each polynomial are $p_1, \ldots, p_n$. In the echelon form the row degrees are the Kronecker indices which we label $\kappa_1, \ldots, \kappa_n$.

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[5]This quotation refers to the echelon form as being complicated, a criticism sometimes made. However, Athanasopoulos, Poskitt and Vahid (2012) relate the echelon form to scalar component models and (page 62) provide "an intuition behind the complicated Echelon form formulae [which] ... demystifies the Echelon form."

We specify a distinction between row degrees $(p_1, \ldots, p_n)$ and Kronecker indices $(\kappa_1, \ldots, \kappa_n)$, even though these are equivalent in the echelon form, since this plays a role in our MCMC algorithm. In this, at one stage we work with a VARMA that simply has row degrees $p_1, \ldots, p_n$, but is otherwise unrestricted. That is, it does not impose the additional restrictions (defined through p_{ik}) required to put the VARMA in echelon form.

As an example of the echelon form, consider a bivariate VARMA$(1, 1)$, denoted as

$$
\begin{pmatrix} 1 & 0 \\ B_{0,21} & 1 \end{pmatrix} \begin{pmatrix} y_{1,t} \\ y_{2,t} \end{pmatrix} = \begin{pmatrix} B_{1,11} & B_{1,12} \\ B_{1,21} & B_{1,22} \end{pmatrix} \begin{pmatrix} y_{1,t-1} \\ y_{2,t-1} \end{pmatrix}
$$
$$
+ \begin{pmatrix} \Theta_{1,11} & \Theta_{1,12} \\ \Theta_{1,21} & \Theta_{1,22} \end{pmatrix} \begin{pmatrix} \epsilon_{1,t-1} \\ \epsilon_{2,t-1} \end{pmatrix} + \begin{pmatrix} 1 & 0 \\ \Theta_{0,21} & 1 \end{pmatrix} \begin{pmatrix} \epsilon_{1,t} \\ \epsilon_{2,t} \end{pmatrix}. \tag{3}
$$

If it is known that $B_{1,21} = B_{1,22} = \Theta_{1,21} = \Theta_{1,22} = 0$, then the conventional restriction $B_{0,21} = \Theta_{0,21} = 0$ is not enough to identify the model. That is, it yields $y_{2,t} = \epsilon_{2,t}$, but in the equation for $y_{1,t}$, $B_{1,12}$ is not separately identified from $\Theta_{1,12}$. To achieve identification in this case, it is further necessary to restrict either $B_{1,12} = 0$ or $\Theta_{1,12} = 0$. However, knowing $B_{1,21} = B_{1,22} = \Theta_{1,21} = \Theta_{1,22} = 0$ implies that the Kronecker indices of the system are $\kappa_1 = 1, \kappa_2 = 0$. Converting (3) to a VARMA$_E(1, 0)$ yields an identified model

$$
\begin{pmatrix} 1 & 0 \\ B_{0,21} & 1 \end{pmatrix} \begin{pmatrix} y_{1,t} \\ y_{2,t} \end{pmatrix} = \begin{pmatrix} B_{1,11} & 0 \\ 0 & 0 \end{pmatrix} \begin{pmatrix} y_{1,t-1} \\ y_{2,t-1} \end{pmatrix}
$$
$$
+ \begin{pmatrix} \Theta_{1,11} & \Theta_{1,12} \\ 0 & 0 \end{pmatrix} \begin{pmatrix} \epsilon_{1,t-1} \\ \epsilon_{2,t-1} \end{pmatrix} + \begin{pmatrix} 1 & 0 \\ B_{0,21} & 1 \end{pmatrix} \begin{pmatrix} \epsilon_{1,t} \\ \epsilon_{2,t} \end{pmatrix}.
$$

Therefore, the rules associated with the echelon form automatically impose the identifying restrictions $B_{1,12} = 0$ and $\Theta_{0,21} = B_{0,21}$, but leave $B_{0,21}$ as a free parameter in the model.

The peculiar aspect of VARMA systems is that if, instead of assuming $B_{1,21} = B_{1,22} = \Theta_{1,21} = \Theta_{1,22} = 0$, we assume that any one of $B_{1,21}$, $B_{1,22}$, $\Theta_{1,21}$, or $\Theta_{1,22}$ is not zero, then the entire system is identified by imposing only the restriction $B_{0,21} = \Theta_{0,21} = 0$. In terms of the echelon form, this corresponds to $\kappa_1 = \kappa_2 = 1$. In general, whenever $\kappa_1 = \cdots = \kappa_n$ in a VARMA$_E$, the model ends up being a conventional unrestricted VARMA (i.e. the semi-structural VARMA with the conventional $B_0 = \Theta_0 = I$ restriction imposed).

In consequence, the key challenge of applying the echelon form methodology in practice is specifying $\boldsymbol{\kappa}$. The problem is that whenever a particular κ_i is over-specified, the resulting VARMA$_E$ is unidentified; whenever it is under-specified, the VARMA$_E$ is misspecified. Therefore, to exploit the theoretical advantages that the VARMA$_E$ provides, the practitioner must choose the Kronecker indices correctly.

The standard frequentist approach to specifying and estimating VARMA models, in consequence, can be described as consisting of three steps:

1. estimate the Kronecker indices, $\hat{\boldsymbol{\kappa}}$;

2. estimate model parameters of the VARMA$_E(\hat{\boldsymbol{\kappa}})$;

3. reduce the model (e.g. using hypothesis testing procedures to eliminate insignificant parameters).

It is important to emphasize that the order of the above steps is crucial. Specifically, step 2 cannot be reasonably performed without completing step 1 first. To appreciate the difficulties with implementing step 1, however, consider performing a full search procedure over all possible Kronecker indices for an n-dimensional system. This would require setting a maximum order $\kappa_{\max}$, estimating $(\kappa_{\max} + 1)^n$ echelon form models implied by each combination of Kronecker indices and then applying some model selection criterion to select the optimal $\boldsymbol{\kappa}$. Given the difficulties associated with maximizing a VARMA$_E$ likelihood, even conditional on a perfectly specified $\boldsymbol{\kappa}$, one cannot hope to complete such a search in a reasonable amount of time (i.e. even a small system with $n = 3$ and $\kappa_{\max} = 5$ would require 1024 Full Information Maximum Likelihood (FIML) routines). Moreover, many of the combinations of $\kappa_1, \ldots, \kappa_n$ that a full search algorithm would need to traverse inevitably result in unidentified specifications, thus plaguing the procedure with exactly the problem that it is designed to resolve.

To handle this difficulty, abbreviated search algorithms relying on approximations are typically employed. Poskitt (1992) provides one particularly popular approach. First, it takes advantage of some special features that arise if the Kronecker indices are re-ordered from smallest to largest such that the number of model evaluations is greatly reduced. Second, it involves a much simpler estimation routine for each evaluation step—i.e., a closed form procedure for consistently (though less efficiently than FIML) estimating the free parameters of a VARMA$_E(\boldsymbol{\kappa})$. These two features also alleviate (although do not eliminate) the problem of needing to estimate unidentified specifications over the course of the search. As a result, consistent estimates of the Kronecker indices are obtained.

However, the implementation also relies on a number of approximations. First, like all existing Kronecker search algorithms, Poskitt (1992) begins by estimating residuals from a long VAR. These are then treated as observations in subsequent least squares estimation routines, which are used to compute information criteria for models of alternative Kronecker structures. Based on the model comparison, the search algorithm terminates when a local optimum is reached. In small samples, therefore, the efficiency of this approach will depend on a number of manual settings and may often lead to convergence difficulties in the likelihood maximization routines implemented at the second stage (for further discussion, see Lutkepohl and Poskitt, 1996).

Consequently, the procedure does not really overcome the basic hurdle: if the $\hat{\boldsymbol{\kappa}}$ obtained in small samples incorrectly describes the underlying structure of the Kronecker indices (as reliable as it may be asymptotically), the VARMA$_E(\hat{\boldsymbol{\kappa}})$ specified in step 2 may ultimately be of little use in resolving the specification and identification issues associated with the unrestricted VARMA.

Recently, Dias and Kapetanios (2013) have developed a computationally-simpler iterated ordinary least squares (OLS) estimation procedure for estimating VARMAs. They prove its consistency and, although it is less efficient than the maximum likelihood estimator (MLE), it has the advantage that it works in places where the MLE does not. In fact, the authors conclude (page 22) that "the constrained MLE algorithm is not a feasible alternative for medium and large datasets due to its computational demand." For instance, they report that their Monte Carlo study which involved 200 artificial generated data sets of 200 observations each from an 8 dimensional VARMA took almost one month of computer time. Their iterated OLS procedure is an approximate method, but

the authors show its potential to work with larger VARMAs such as those considered in the present paper. However, their method does run into the problem that it can often fail to converge when either the sample size is small or the dimension of the VARMA is large. For instance, their various Monte Carlo exercises report failure to convergence rates from 79% to 97% for VARMAs with 10 dependent variables and $T = 150$. These results are generated with VARMA(1,1) models and would, no doubt, worsen with longer lag lengths such as those we use in the present paper. These high failure to converge rates are likely due to the fact that, with many parameters to estimate and relatively little data to inform such estimates, likelihood functions (or approximations to them) can be quite flat and their optima difficult to find. This motivates one theme of our paper: use of carefully selected shrinkage through a Bayesian prior is useful in producing sensible (and computationally feasible) results in large VARMA models.

It is not difficult to see why applied macroeconomists have rarely used these frequentist procedures for estimating VARMAs. To extend them to models with stochastic volatility or a similar feature commonly incorporated in modern VARs seems extremely difficult. Shortly, we will develop a Bayesian method which jointly goes through the three steps listed above in the context of a single MCMC algorithm and allows for many extensions (e.g. adding stochastic volatility) in a straightforward fashion. Before we do this, we describe an alternative way of parameterizing the VARMA which is used in our MCMC algorithm.

2.3 The Expanded Form for the VARMA

Papers such as Metaxoglou and Smith (2007) and Chan and Eisenstat (2015) adopt an alternative way of parameterizing the VARMA called the expanded VARMA form which proves useful for computational purposes. The expanded VARMA form can be written as:

$$\mathbf{B}_0 \mathbf{y}_t = \sum_{j=1}^{p} \mathbf{B}_j \mathbf{y}_{t-j} + \sum_{j=0}^{p} \mathbf{\Phi}_j \mathbf{f}_{t-j} + \boldsymbol{\eta}_t, \tag{4}$$

where $\mathbf{f}_t \sim \mathcal{N}(\mathbf{0}, \mathbf{\Omega})$ and $\boldsymbol{\eta}_t \sim \mathcal{N}(\mathbf{0}, \mathbf{\Lambda})$ are independent, $\mathbf{\Phi}_0$ is a lower triangular matrix with ones on the diagonal, $\mathbf{\Phi}_1, \ldots, \mathbf{\Phi}_p$ are coefficient matrices, and $\mathbf{\Omega}, \mathbf{\Lambda}$ are diagonal.

As discussed in Metaxoglou and Smith (2007) and Chan and Eisenstat (2015), the expanded form is an equivalent representation of the VARMA in (1), albeit an over-parameterized one.[6] The over-parametrization can be regarded as arising from the additional n parameters $\mathbf{\Lambda}_{11}, \ldots, \mathbf{\Lambda}_{nn}$. However, given the parameters $\mathbf{B}_0, \mathbf{B}_1, \ldots, \mathbf{B}_p$, $\mathbf{\Phi}_0, \mathbf{\Phi}_1, \ldots, \mathbf{\Phi}_p, \mathbf{\Omega}$, and $\mathbf{\Lambda}$ of the expanded form, the VARMA parameters $\mathbf{B}_0, \mathbf{B}_1, \ldots, \mathbf{B}_p$, $\mathbf{\Theta}_1, \ldots, \mathbf{\Theta}_p$, and $\mathbf{\Sigma}$ can be easily computed because the mapping from the expanded form parameters to the semi-structural VARMA parameters is always well defined. Specifically, $\mathbf{B}_0, \ldots, \mathbf{B}_p$ are equivalent in both representations and $\mathbf{\Theta}_1, \ldots, \mathbf{\Theta}_p, \mathbf{\Sigma}$ can be recovered using the procedure developed in Chan and Eisenstat (2015), which is reproduced for convenience in our online appendix.

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[6]The theoretical foundation for this statement is found in Peiris (1988), Theorem 2, which states that the sum of any two VMA processes results in a VMA process, and therefore, any VMA process can be decomposed into a VMA plus white noise.

In light of this, Chan and Eisenstat (2015) propose building an MCMC sampling scheme directly on the expanded form in (4) and recovering draws of the VARMA parameters in (1) by applying the previously discussed transformation on each draw of the expanded form parameters. In doing so, it is important to emphasize that while $\boldsymbol{\Phi}_0, \boldsymbol{\Phi}_1, \ldots, \boldsymbol{\Phi}_p, \boldsymbol{\Omega}$, and $\boldsymbol{\Lambda}$ are not identified point-wise (note, however, that $\mathbf{B}_0, \ldots, \mathbf{B}_p$ are nevertheless identified in the expanded form), invertibility of $\boldsymbol{\Theta}(L)$ and positive-definiteness of $\boldsymbol{\Sigma}$ imply substantial restrictions on the expanded form parameters. Chan and Eisenstat (2015) examine the theoretical properties of the mapping between the expanded form and the semi-structural VARMA form and demonstrate that, in fact, Λ_{ii} are always identified up to a finite interval. This interval is largest as the $\boldsymbol{\Theta}_j$ tend to zero and give rise to the restriction that $\boldsymbol{\Sigma} - \boldsymbol{\Lambda}$ is p.s.d.; the interval on all Λ_{ii} collapses to zero as all roots of $\boldsymbol{\Theta}(L)$ approach the unit circle.

In consequence, working with the expanded form does not require restrictive priors on $\boldsymbol{\Phi}_0, \boldsymbol{\Phi}_1, \ldots, \boldsymbol{\Phi}_p, \boldsymbol{\Omega}, \boldsymbol{\Lambda}$ and we follow Chan and Eisenstat (2015) in specifying default, weakly informative priors on these parameters (with extension to shrinkage on $\boldsymbol{\Phi}_j$), as discussed extensively in subsequent sections and the online appendix. At the same time, the expanded form is a linear state space model, and therefore, admits a straightforward and efficient Gibbs sampling algorithm. Moreover, one does not need to impose invertibility restrictions directly at each iteration of the MCMC algorithm as these can be easily applied in the *ex post* processing of the draws (see the online appendix and Chan and Eisenstat, 2015, for more details). Chan and Eisenstat (2015) provide details of how such an algorithm may be efficiently implemented in simple VARMA settings. However, they do not consider canonical VARMAs and the implementation of echelon form restrictions.

In this paper we build on the computational advantages offered by the expanded form and develop Gibbs sampling algorithms that focus on the canonical echelon specification and parsimony in large system where shrinkage is indispensable. One key extension, in this respect, is that our new Gibbs sampler features a stochastic search for echelon form structures. This is possible owing to the fact that all echelon restrictions on $\boldsymbol{\Theta}_j$ for $1 \leq j \leq p$ (see subsection 2.2) will correspond to equivalent restrictions on $\boldsymbol{\Phi}_j$. In consequence, imposing echelon form restrictions on $\mathbf{B}_0, \ldots, \mathbf{B}_p, \boldsymbol{\Phi}_1, \ldots, \boldsymbol{\Phi}_p$ in the course of the Gibbs sampler will generate equivalent echelon form restrictions on the $\mathrm{VARMA}_E(\boldsymbol{\kappa})$ recovered *ex post*.[7] At the conclusion of this procedure, we obtain draws of $\mathbf{B}_0, \ldots, \mathbf{B}_p, \boldsymbol{\Theta}_0, \ldots, \boldsymbol{\Theta}_p$ and Σ that always satisfy echelon form restrictions conditional on the corresponding draw of $\boldsymbol{\kappa}$. Inference is then based on draws of these parameters.

3 Bayesian Inference in VARMA Models

3.1 The Existing Bayesian Literature

Previously, we have drawn a distinction between the related concepts of parsimony and identification. Identification can be achieved by selecting the correct Kronecker indices (which imply certain restrictions on a semi-structural VARMA model). Parsimony is a

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[7]This preservation of echelon form restrictions is not affected by the imposition of invertibility in the post-processing of draws either.

more general concept, involving either setting coefficients to zero (or any constant) or shrinking them towards zero. So identification can be achieved through parsimony (i.e. selecting the precise restrictions implied by the Kronecker indices in the context of an unidentified VARMA model), but parsimony can involve imposing other restrictions on a non-identified model or imposing restrictions beyond that required for identifying the model.

In this sense, the Bayesian literature breaks into two groups. The first consists of papers which estimate VARMA models, possibly taking into account parsimony considerations. Good examples of this literature are Ravishanker and Ray (1997) and Chan and Eisenstat (2015). The second consists of papers which explicitly address identification issues. The key references in this strand of the literature is Li and Tsay (1998). Since one important aspect of our paper lies in identification, we will focus on this paper.

Li and Tsay (1998) specify a model similar to (1) but parameterized somewhat differently (i.e. their $\boldsymbol{\Theta}_0$ is lower triangular but not equal to $\mathbf{B}_0$ and, thus, they work with a diagonal $\boldsymbol{\Sigma}$) and work with the echelon form, attempting to jointly estimate the VARMA parameters with the Kronecker indices. This is done through the use of a hierarchical prior for the coefficients which is often called a stochastic search variable selection (SSVS) prior (although other terminologies exist). Before describing Li and Tsay's algorithm, we briefly introduce the idea underlying SSVS in a generic context. Let α be a parameter. SSVS specifies a hierarchical prior (i.e. a prior expressed in terms of parameters which in turn have a prior of their own) which is a mixture of two Normal distributions:

$$\alpha \mid \gamma \sim (1 - \gamma)\mathcal{N}(0, \tau_0^2) + \gamma\mathcal{N}(0, \tau_1^2), \tag{5}$$

where $\gamma \in \{0, 1\}$ we refer to as an indicator variable. Thus, if $\gamma = 1$ then the prior for α is given by the second Normal and if $\gamma = 0$ it is given by the first Normal. The prior is hierarchical since γ is treated as an unknown parameter and estimated in a data-based fashion. The aspect which allows for prior shrinkage and variable selection arises by choosing the first prior variance, τ_0^2, to be "small" (so that the coefficient is shrunk so as to be close to zero) and the second prior variance, τ_1^2, to be "large" (implying a relatively noninformative prior for the corresponding coefficient). An SSVS prior of this sort, which we shall call "soft SSVS", has been used by many researchers. For instance, George, Sun and Ni (2008) and Koop (2013) use it with VARs and Li and Tsay (1998) adopt something similar. An extreme case of the SSVS prior arises if the first Normal in (5) is replaced by a point mass at zero. This we will call "hard SSVS". It was introduced in Kuo and Mallick (1997) and used with VARs by Korobilis (2013) and others.

Li and Tsay (1998) specify soft SSVS priors on the VAR and VMA coefficients of a VARMA. The ingenuity of this approach is that it combines in practical terms the two related concepts of identification and parsimony. The authors enforce the echelon form through this framework by imposing certain deterministic relationships between the SSVS indicators (see section 4 of Li and Tsay, 1998, for more details). Based on this, they devise an MCMC algorithm that cycles through n individual (univariate) ARMAX equations. The ith ARMAX equation is obtained by treating the observations $\{y_{k,t}\}$ for $k = 1, \ldots, i - 1$, $t = 1, \ldots, T$ and the computed errors $\{\epsilon_{k,t}\}$ for $k \neq i, t = 1, \ldots, T$ as exogenous regressors. SVSS indicators are then updated conditional on draws of the

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coefficients and subject to the deterministic relationships implied by the echelon form. In consequence, draws of the Kronecker indices (which can be recovered from draws of the SSVS indicators) are simultaneously generated along with the model parameters.

Their algorithm, however, entails a significant degree of complexity both in terms of programming and computation. A pass through each equation requires reconstructing VARMA errors (i.e. based on previous draws of parameters pertaining to other equations) and sampling three parameter blocks: (i) the autoregressive and "exogenous" variable coefficients, (ii) the error variance, and (iii) the moving average parameters. The latter entails a non-trivial Metropolis-Hastings step, and all must be repeated n times for every sweep of the MCMC routine. Evidently, the complexity of this algorithm grows rather quickly with the size of the system, and in their applications, only systems with $n = 3$ and $\kappa_{\max} \leq 3$ are considered. The run times reported for even these small systems are measured in hours.

Relative to Li and Tsay (1998) our algorithm shares the advantage of jointly estimating Kronecker indices and model parameters, thus ensuring parsimony and identification. However, we argue that ours is a more natural specification, which also provides great computational benefits and allows us to work with the large Bayesian VARMAs of interest to empirical macroeconomists, whereas the Li and Tsay (1998) algorithm is computationally infeasible in large dimensional settings. First, by using the expanded form discussed in subsection 2.3, we are able to work with a familiar, linear state space model. Conditional on the Kronecker indices, computation is fast and efficient even for large n. Moreover, this representation enables us to analytically integrate out the coefficients $\{\mathbf{B}_j\}$ and $\{\mathbf{\Phi}_j\}$ when sampling the Kronecker indices. The efficiency gains from this are particularly important as n increases because the size of each $\mathbf{B}_j$ and $\mathbf{\Phi}_j$ grows quadratically with n. In fact, this added efficiency together with the reduced computational burden is precisely what allows us to estimate an exact echelon form VARMA for large systems. The details are provided in the following subsection.

3.2 Our Approach to Bayesian Inference in VARMAs

Our approach to Bayesian inference is based on the ideas that identification is achieved in the echelon form (i.e. through estimating $\boldsymbol{\kappa}$ in the VARMA$_E(\boldsymbol{\kappa})$), but computation is more easily done in the expanded form (see also Chan and Eisenstat, 2015). Thus, our MCMC algorithm works in the latter, but draws are transformed to the echelon form *ex post*, and inference is drawn from the model in (1). We also treat the Kronecker indices $\boldsymbol{\kappa}$ as unknown and sample them with a stochastic search algorithm. Parsimony and identification are achieved using SSVS priors.

Remember that we have three kinds of restrictions which may be of interest in the echelon form VARMA. First, a given $\boldsymbol{\kappa}$ implies restrictions on the row degrees of each equation. Second, the value for $\boldsymbol{\kappa}$ implies additional restrictions on the VARMA coefficients (see Section 2.2 for the definition of the echelon form and a reminder of what these restrictions are). Third, we may have other restrictions which have nothing to do with the identification restrictions implied by $\boldsymbol{\kappa}$, but may be worth imposing solely for reasons of parsimony. We use a hierarchical SSVS prior that automatically allows for the imposition (or not) of each of these types of restriction. We outline two SSVS priors which differ in

their treatment of the second set of restrictions. Our main SSVS prior always imposes the echelon form implied by $\boldsymbol{\kappa}$. That is, its main component imposes the first two sets of restrictions. In this case, draws of $p_1, \ldots, p_n$ are equivalent to draws of $\kappa_1, \ldots, \kappa_n$ (and, hence, we can parameterize in terms of either and we choose $p_1, \ldots, p_n$ below). However, as we shall see, allowing for the imposition of the two sets of restrictions implied by the echelon form introduces dependencies in the hierarchical prior which slow down computation. Hence, we also introduce an alternative SSVS prior whose main component only imposes (or not) the first set of restrictions. Thus, the main part of this alternative prior can impose some, but not all, of the restrictions implied by the echelon form. The remaining part of our hierarchical prior is a conventional SSVS prior which can impose restrictions on any individual coefficient. In the main algorithm, this conventional SSVS prior can be used to impose parsimony beyond that required for identification. In the alternative algorithm, this conventional SSVS prior is used for both the second and third set of restrictions. Thus, in this alternative algorithm, it is possible (but not necessary) for this additional SSVS prior to impose the identifying restrictions missed by the main part of the prior. The remainder of this section provides details of how this is done.

Since row degree restrictions are especially important for identifying the lag structure, we always use hard SSVS for these (i.e., the restrictions implied by a particular choice of $p_1, \ldots, p_n$ are imposed exactly). Restrictions on the remaining parameters are partly used for identification (i.e. when the echelon is enforced exactly, addition restrictions beyond those implied solely by a choice for $p_1, \ldots, p_n$ are required) and partly to achieve additional parsimony (i.e., by further restricting parameters which remain in the $\text{VARMA}_E(\boldsymbol{\kappa})$). For these, the researcher may wish to use either soft or hard SSVS and, in this paper, we allow for both. With some abuse of terminology, we will call the prior which uses hard SSVS to achieve identification and soft SSVS to achieve parsimony the "soft SSVS prior" and the prior which imposes hard SSVS throughout the "hard SSVS prior". In what follows, we describe the main features of our approach, paying particular attention to the SSVS priors on the VARMA coefficients. Complete details on the priors for the remaining parameters are given in the online appendix.

Consider the expanded form VARMA given in (4) for which the VARMA coefficients are parameterized in terms of $\mathbf{B}_j$ and $\boldsymbol{\Phi}_j$. Let the individual coefficients in these matrices be denoted $B_{j,ik}$ and $\Phi_{j,ik}$, respectively. Here we describe the soft SSVS implementation with $\tau^2_{0,j,ik} \ll \tau^2_{1,j,ik}$ (the hard SSVS implementation will be the same except there is no $\tau^2_{0,j,ik}$, but instead a point mass at zero is used) which is given by

$$
\begin{aligned}
\left(B_{j,ik} \,|\, p_i, \gamma^{B,S}_{j,ik} \right) &\sim \left(1 - \gamma^{B,R}_{j,ik} \right) \mathbb{1}(B_{j,ik} = 0) \\
&\quad + \gamma^{B,R}_{j,ik} \left(\left(1 - \gamma^{B,S}_{j,ik} \right) \mathcal{N}(0, \tau^2_{0,j,ik}) + \gamma^{B,S}_{j,ik} \mathcal{N}(0, \tau^2_{1,j,ik}) \right), \\
\left(\Phi_{j,ik} \,|\, p_i, \gamma^{\Phi,S}_{j,ik} \right) &\sim \left(1 - \gamma^{\Phi,R}_{j,ik} \right) \mathbb{1}(\Phi_{j,ik} = 0) \\
&\quad + \gamma^{\Phi,R}_{j,ik} \left(\left(1 - \gamma^{\Phi,S}_{j,ik} \right) \mathcal{N}(0, \tau^2_{0,j,ik}) + \gamma^{\Phi,S}_{j,ik} \mathcal{N}(0, \tau^2_{1,j,ik}) \right),
\end{aligned}
\tag{6}
$$

where $\mathbb{1}(\cdot)$ is the indicator function. In this setup, $\gamma^{B,R}_{j,ik}, \gamma^{\Phi,R}_{j,ik} \in \{0,1\}$ are indicators used solely to impose restrictions on the row degrees (R denoting "row degree"): $\gamma^{B,R}_{j,ik} = \gamma^{\Phi,R}_{j,ik} =$

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1 iff $0 < j \leq p_i$ or $j = 0, i < k$. In words, p_i is the row degree of equation i, and, thus, all the coefficients with lag length greater than p_i are set to zero. This is obtained by setting $\gamma_{j,ik}^{B,R} = \gamma_{j,ik}^{\Phi,R} = 0$ for lag lengths longer than p_i (i.e. $j > p_i$). A given value of p_i tells us exactly what values $\gamma_{j,ik}^{B,R}, \gamma_{j,ik}^{\Phi,R}$ take in equation i. This justifies how we can treat $p_1, \ldots, p_n$ as the model parameters (which we sample directly) and $\{\gamma_{j,ik}^{B,R}, \gamma_{j,ik}^{\Phi,R}\}$ as transformations of $p_1, \ldots, p_n$ (i.e. through the mapping $\gamma^R = \{\gamma_{j,ik}^{B,R}, \gamma_{j,ik}^{\Phi,R}\} = \mathcal{R}(p_1, \ldots, p_n)$). Furthermore, $\gamma_{j,ik}^{B,S}, \gamma_{j,ik}^{\Phi,S} \in \{0, 1\}$ are the indicators related to the SSVS mechanism for the remaining coefficients not restricted by the row degrees (S denoting "shrinkage").

This prior is applied to the expanded VARMA. In order to impose the identifying restrictions implied by $\boldsymbol{\kappa}$ on the echelon form, we need to both restrict the row degrees appropriately (using $\gamma_{j,ik}^{B,R}, \gamma_{j,ik}^{\Phi,R}$) and impose the additional restrictions needed to define the echelon form (using $\gamma_{j,ik}^{B,S}, \gamma_{j,ik}^{\Phi,S}$). To see how this works, define a new set of echelon form indicators and a mapping from row degrees (or, equivalently, Kronecker indices) to the indicators: $\boldsymbol{\gamma}^E = \{\gamma_{j,ik}^{B,E}, \gamma_{j,ik}^{\Phi,E}\} = \mathcal{E}(p_1, \ldots, p_n)$. The echelon form can be imposed using γ^R and prior on $\gamma_{j,ik}^{B,S}$ conditional on $p_1, \ldots, p_n$ of the form

$$\Pr\left(\gamma_{j,ik}^{B,S} = 1 \mid p_1, \ldots, p_n\right) = \begin{cases} 0 & \text{if } \gamma_{j,ik}^{B,E} = 0 \text{ and } \gamma_{j,ik}^{B,R} = 1 \\ 0.5 & \text{otherwise} \end{cases}. \tag{7}$$

To further clarify this setup, recall the bivariate VARMA(1,1) example of subsection 2.2 which involves two row degrees, p_1 and p_2. Table 1 depicts the mapping $\{\gamma_{j,ik}^{B,E}, \gamma_{j,ik}^{\Phi,E}\} = \mathcal{E}(p_1, p_2)$ and Table 2 depicts the mapping $\{\gamma_{j,ik}^{B,R}, \gamma_{j,ik}^{\Phi,R}\} = \mathcal{R}(p_1, p_2)$. Observe that for any p_1, p_2, the set of zero restrictions prescribed by $\mathcal{R}(p_1, p_2)$ is always a subset of the zero restrictions prescribed by $\mathcal{E}(p_1, p_2)$. For example, when $p_1 = 0, p_2 = 1$ both $R(p_1, p_2)$ and $\mathcal{E}(p_1, p_2)$ lead to $B_{1,11} = B_{1,12} = \Phi_{1,11} = \Phi_{1,12} = 0$ and $B_{1,22}, \Phi_{1,21}, \Phi_{1,22}$ unrestricted. However, $\mathcal{E}(p_1, p_2)$ further imposes the two addition restrictions $B_{0,21} = B_{1,21} = 0$ (i.e. $\gamma_{0,21}^{B,E} = \gamma_{1,21}^{B,E} = 0$) while $\mathcal{R}(p_1, p_2)$ does not (i.e. $\gamma_{0,21}^{B,R} = \gamma_{1,21}^{B,R} = 1$).

To guarantee that the ensuing MCMC algorithm always generates draws from an exact echelon form VARMA, the prior in (7) relegates the two additional restrictions imposed by $\mathcal{E}(p_1, p_2)$ (but not $\mathcal{R}(p_1, p_2)$) to the SSVS indicators, as depicted in Table 3.

Note that this construction has two important implications. First, the additional restrictions will be soft whenever the soft SSVS prior is used (i.e. $\tau_{0,j,ik}^2$ is small) and hard only when the hard SSVS prior is used (i.e. $\tau_{0,j,ik}^2 = 0$). Using the soft SSVS prior in the example above, consequently, would lead to $B_{0,21}$ and $B_{1,21}$ being sampled with a small variance (conditional on $p_1 = 0, p_2 = 1$) and $B_{1,11}, B_{1,12}, \Phi_{1,11}, \Phi_{1,12}$ being set to zero with probability 1. However, the fact that a part of the echelon form restrictions are implemented as soft restrictions is of little empirical consequence. Of greater importance are the restrictions implied by row degrees since these define the lag structure of each equation, and we prefer to impose these in exact fashion.

Second, we must have $\Pr\left(\gamma_{j,ik}^{B,S} = 1 \mid p_1, \ldots, p_n\right) \neq 1$ whenever $p_1, \ldots, p_n$ implies $\gamma_{j,ik}^{B,E} = 1$ or $\gamma_{j,ik}^{B,R} = 0$. Otherwise, the Gibbs sampler constructed below would yield a reducible Markov chain (we return to this point in subsection 3.3 below). The important implication is that shrinkage priors on all coefficients (including those not restricted by identification) form an integral part of the stochastic search algorithm. Setting

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$\Pr\left(\gamma_{j,ik}^{B,S} = 1 \,|\, p_1, \ldots, p_n\right) = 0.5$ (whenever $\gamma_{j,ik}^{B,E} = 1$ or $\gamma_{j,ik}^{B,R} = 0$) is desirable since the value 0.5 implies a restriction is, a priori, equally likely to apply as not; hence, it provides the most flexibility for the algorithm to uncover relevant echelon form structures.

Table 1: Echelon form indicators $\{\gamma_{j,ik}^{B,E}, \gamma_{j,ik}^{\Phi,E}\}$ in the bivariate VARMA$(1,1)$ example.

p_1	p_2	$\gamma_{0,21}^{B,E}$	$\gamma_{1,11}^{B,E}$	$\gamma_{1,21}^{B,E}$	$\gamma_{1,12}^{B,E}$	$\gamma_{1,22}^{B,E}$	$\gamma_{1,11}^{\Phi,E}$	$\gamma_{1,21}^{\Phi,E}$	$\gamma_{1,12}^{\Phi,E}$	$\gamma_{1,22}^{\Phi,E}$
0	0	0	0	0	0	0	0	0	0	0
0	1	0	0	0	0	1	0	1	0	1
1	0	1	1	0	0	0	1	0	1	0
1	1	0	1	1	1	1	1	1	1	1

Table 2: Row degree indicators $\{\gamma_{j,ik}^{B,R}, \gamma_{j,ik}^{\Phi,R}\}$ in the bivariate VARMA$(1,1)$ example.

p_1	p_2	$\gamma_{0,21}^{B,R}$	$\gamma_{1,11}^{B,R}$	$\gamma_{1,21}^{B,R}$	$\gamma_{1,12}^{B,R}$	$\gamma_{1,22}^{B,R}$	$\gamma_{1,11}^{\Phi,R}$	$\gamma_{1,21}^{\Phi,R}$	$\gamma_{1,12}^{\Phi,R}$	$\gamma_{1,22}^{\Phi,R}$
0	0	0	0	0	0	0	0	0	0	0
0	1	1	0	1	0	1	0	1	0	1
1	0	1	1	0	1	0	1	0	1	0
1	1	1	1	1	1	1	1	1	1	1

Table 3: SSVS prior $\Pr\left(\gamma_{j,ik}^{B,S} = 1 \,|\, p_1, \ldots, p_n\right)$ for imposing the echelon form in the bivariate VARMA$(1,1)$ example.

p_1	p_2	$\gamma_{0,21}^{B,S}$	$\gamma_{1,11}^{B,S}$	$\gamma_{1,21}^{B,S}$	$\gamma_{1,12}^{B,S}$	$\gamma_{1,22}^{B,S}$	$\gamma_{1,11}^{\Phi,S}$	$\gamma_{1,21}^{\Phi,S}$	$\gamma_{1,12}^{\Phi,S}$	$\gamma_{1,22}^{\Phi,S}$
0	0	0	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5
0	1	0	0.5	0	0.5	0.5	0.5	0.5	0.5	0.5
1	0	0.5	0.5	0.5	0	0.5	0.5	0.5	0.5	0.5
1	1	0	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5

To complete the prior specification, we set $\Pr\left(\gamma_{j,ik}^{\Phi,S} = 1\right) = 0.5$ for the indicators on the elements of $\mathbf{\Phi}_j$ and uniform priors on $p_1, \ldots, p_n$, which induce a prior on $\boldsymbol{\gamma}^R$, and by implication, a uniform prior on the Kronecker indices.[8] Our MCMC algorithms provide draws of $p_1, \ldots, p_n$, and under the prior specification (7), these are equivalent to draws of the Kronecker indices $\kappa_1, \ldots, \kappa_n$. Parameters of interest in terms of (1) can be recovered from draws of $\mathbf{B}_0, \mathbf{B}_1, \ldots, \mathbf{B}_p, \mathbf{\Phi}_0, \mathbf{\Phi}_1, \ldots, \mathbf{\Phi}_p, \mathbf{\Omega}$, and $\mathbf{\Lambda}$ using the procedure described in the online appendix. As point out in subsection 2.3, the echelon form is preserved under the latter transformation.

In this framework, a particular identification scheme can be imposed through a dogmatic prior which sets probability one to a particular value for κ (e.g. allocating prior

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[8]Since $\mathcal{R}(p_1, ..., p_n)$ and $\mathcal{E}(p_1, ..., p_n)$ always yield $\gamma_{j,ik}^{\Phi,R} = \gamma_{j,ik}^{\Phi,E}$, priors on $\gamma_{j,ik}^{\Phi,S}$ are not restricted by echelon form considerations and other priors can easily be accommodated.

probability one to $\kappa_1 = \cdots = \kappa_n = p$ will be equivalent to estimating an unrestricted VARMA(p,p)). In this case, we can work directly with $\boldsymbol{\gamma}^E$ (i.e. instead of $\boldsymbol{\gamma}^R$) to enforce the echelon form restrictions, and the SSVS indicators $\boldsymbol{\gamma}^S = \{\gamma_{j,ik}^{B,S}, \gamma_{j,ik}^{\Phi,S}\}$ would then be used exclusively to control additional shrinkage: they can either be fixed *a priori* with $\mathbb{P}(\gamma_{j,ik}^{\cdot,S} = 1) = 1$ such that no additional shrinkage/variable selection is employed, or specified as $\mathbb{P}(\gamma_{j,ik}^{\cdot,S} = 1) = 0.5$ and sampled in the course of the MCMC run along with the other parameters. Applying the latter and naively setting $\kappa_1 = \cdots = \kappa_n = p$ leads to a simple SSVS model where the parameters are potentially unidentified, but parsimony is achieved through shrinkage and computation is very fast. In either case, treating $\boldsymbol{\kappa}$ as fixed eliminates the need to use SSVS indicators for enforcing the echelon form.

Working with stochastic $\boldsymbol{\kappa}$ through stochastic row degrees $p_1, \ldots, p_n$ and indicators $\gamma_{j,ik}^{B,S}$, $\gamma_{j,ik}^{\Phi,S}$ as outlined above, on the other hand, results in an algorithm that always operates on a parameter space restricted according the echelon form, but also allows for additional shrinkage on the unrestricted coefficients. One interesting consequence of this is that, unlike the classic VARMA$_E(\boldsymbol{\kappa})$ model in which the number of AR coefficients must equal the number of MA coefficients, the additional SSVS priors allows the stochastic search algorithm to uncover a VARMA(p,q) where $p \neq q$ (i.e. if the SSVS mechanism additionally forces certain coefficients to zero).

In sum, we argue that this SSVS prior can successfully address two of the three reasons (identification and parsimony) for a dearth of empirical work which uses VARMAs outlined in the introduction. The third reason was computation. Our MCMC algorithm, outlined below and detailed in the online appendix, is fairly efficient and we have had success using it in quite large VARMAs. For instance, we present empirical work below for VARMAs with $n = 12$ which is much larger than anything we have found in the existing literature with the exception of Dias and Kapetanios (2013). However, dealing with much higher dimensional models (e.g. $n = 25$ or more) as has been sometimes done with VARs would represent a serious, possibly insurmountable computational burden, with our algorithm.

For these reasons, we also consider an approximate MCMC algorithm which is much simpler. This latter algorithm is achieved by replacing (7), which involves prior dependencies between restrictions, with the simpler independent choice $\Pr\left(\gamma_{j,ik}^{B,S} = 1\right) = 0.5$. In our artificial data experiments (see the online appendix), this approximate algorithm (which we call the "row degree algorithm") seems to work quite well and is much more efficient than our exact algorithm (which we call the "echelon algorithm"). Complete details are given in the online appendix, but to understand the intuition underlying the approximate algorithm observe that (7) creates cross-equation relationships among indicators, and therefore, strong dependence between the row degrees $p_1, \ldots, p_n$. For MCMC, this forces us to sample each p_i conditional on all other row degrees and keep track of all these relationships.

However, simplifying the prior on $\gamma_{j,ik}^{B,S}$ as above allows the approximate row degree algorithm to just draw from the row degrees ignoring the other restrictions implied by the echelon form.[9] In this case, the row degrees are conditionally independent of one another

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[9] Note that in this case as with a fixed $\boldsymbol{\kappa}$, SSVS indicators are once again *a priori* separate of any identification restrictions.

and the MCMC algorithm becomes much more efficient. This algorithm has the drawback of ignoring some restrictions of the echelon form, and therefore, the echelon form is not guaranteed to be preserved at every MCMC iteration. Nevertheless this drawback, in practice, may be slight since the SSVS prior on the VARMA coefficients (i.e. involving $\boldsymbol{\gamma}^S$) should be able to pick up any restrictions missed by using an approximate algorithm. Thus, the row degree algorithm may be useful for the researcher who finds our echelon form algorithm too computationally demanding.

3.3 Overview of MCMC Algorithms

The hierarchical prior laid out in the previous subsection, combined with the linear state space nature of the expanded form (4) gives rise to a fairly straightforward Gibbs sampling algorithm. The five major steps can be summarized as follows:

1. Sample $\left(\boldsymbol{p} \,|\, \boldsymbol{\gamma}^S, \mathbf{f}, \boldsymbol{\Lambda}, \mathbf{y}\right)$ *marginal* of $\mathbf{B}, \boldsymbol{\Phi}$, where $\boldsymbol{p} = (p_1, \ldots, p_n)$.

2. Sample $\left(\boldsymbol{\gamma}_i^S, \mathbf{B}_{(i)}, \boldsymbol{\Phi}_{(i)} \,|\, \boldsymbol{p}, \mathbf{f}, \Lambda_{ii}, \mathbf{y}_i\right)$ for each $i = 1, \ldots, n$, where $\mathbf{B}_{(i)}$ denotes the i-th row of $\mathbf{B} = (\mathbf{I}_n - \mathbf{B}_0, \mathbf{B}_1, \ldots, \mathbf{B}_p)$, $\boldsymbol{\Phi}_{(i)}$ the i-th row of $\boldsymbol{\Phi} = (\boldsymbol{\Phi}_0, \ldots, \boldsymbol{\Phi}_p)$, and $\boldsymbol{\gamma}_i^S$ is the set of all SSVS indicators pertaining to $\mathbf{B}_{(i)}, \boldsymbol{\Phi}_{(i)}$.

3. Sample $\left(\Lambda_{ii} \,|\, \mathbf{B}_{(i)}, \boldsymbol{\Phi}_{(i)}, p_i, \boldsymbol{\gamma}_i^S, \mathbf{f}, \mathbf{y}_i\right)$ for each $i = 1, \ldots, n$.

4. Sample $\left(\Omega_{ii} \,|\, \mathbf{f}_i\right)$ for each $i = 1, \ldots, n$.

5. Sample $\left(\mathbf{f} \,|\, \mathbf{B}, \boldsymbol{\Phi}, \boldsymbol{\Omega}, \boldsymbol{\Lambda}, \boldsymbol{p}, \boldsymbol{\gamma}^S, \mathbf{y}\right)$.

Note that Step 1 will also provide draws of γ^R through the mapping $\gamma^R = \mathcal{R}(p_1, \ldots, p_n)$. Steps 1 and 2 above may need to be broken down into further conditional distributions, depending on the particular algorithm under consideration. However, the sampler will always involve drawing from analytically tractable distributions such that no Metropolis-Hastings steps are required. In consequence, the algorithm requires little tuning beyond specifying prior hyperparameters. As discussed in preceding sections and further detailed in the online appendix, we specify standard, weakly informative priors on the parameters of the expanded form. In extensive experimentation with various (stationary and standardized) data sets, we have found the algorithm to perform well using the same default specifications. The subsequent empirical section provides more details on the efficacy and efficiency of the algorithm with macroeconomic data. The online appendix offers additional evidence relating to the algorithms using artificial data.

Specific computational details regarding each Gibbs step are discussed in the online appendix. Here we will highlight the key aspects of Steps 1 and 2 that constitute the sampling of row degrees and SSVS indicators.[10] Related to Step 1, consider first the row degree algorithm where (7) is replaced by the prior $\Pr\left(\gamma_{j,ik}^{B,S} = 1\right)$ that is independent of

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[10]Steps 3-5 involve sampling from Gamma and multivariate Normal distributions.

$p_1, \ldots, p_n$, and therefore,

$$p\left(\boldsymbol{p} \,|\, \boldsymbol{\gamma}^S, \mathbf{f}, \boldsymbol{\Lambda}, \mathbf{y}\right) = \prod_{i=1}^{n} \Pr\left(p_i = l \,|\, \boldsymbol{\gamma}_i^S, \mathbf{f}, \Lambda_{ii}, \mathbf{y}_i\right)$$

$$\Pr\left(p_i = l \,|\, \boldsymbol{\gamma}_i^S, \mathbf{f}, \Lambda_{ii}, \mathbf{y}_i\right) \propto p\left(\mathbf{y}_i \,|\, p_i = l, \boldsymbol{\gamma}_i^S, \mathbf{f}, \Lambda_{ii}\right).$$

Two observations of this sampling step are noteworthy. First, because the expanded form represents a linear state space, we can analytically integrate out the coefficients $\mathbf{B}, \boldsymbol{\Phi}$ conditional on $\mathbf{f}$ such that row degrees are sampled marginally of these coefficients. This is a feature of both the row degree and echelon algorithms, and it gains importance as the size of the model increases because the number of parameters in $\mathbf{B}$ and $\boldsymbol{\Phi}$ grows quadratically with n. Details on how the collapsed likelihood $p\left(\mathbf{y}_i \,|\, p_i = l, \boldsymbol{\gamma}_i^S, \mathbf{f}, \Lambda_{ii}\right)$ is efficiently evaluated are provided in the online appendix.

Second, the row degree algorithm further permits sampling $p_1, \ldots, p_n$ jointly and only requires the partial likelihood $p(\mathbf{y}_i \,|\, \cdot)$ to be evaluated for each possible value of p_i. Both of these features provide significant advantages in larger models as they simultaneously improve sampling efficiency and reduce the computational burden within each Gibbs step.

For the echelon algorithm, a particular set of row degrees implies cross equation restrictions, and therefore, we need to sample p_i conditional on all other row degrees $\boldsymbol{p}_{-i}$. Specifically, for each proposed value of p_i conditional on $\mathbf{p}_{-i}$ and a set of SSVS indicators $\boldsymbol{\gamma}^S$, we need to ensure that a valid echelon form would be preserved. Thus, for each possible value $p_i = l$, it is necessary to compute the implied echelon restrictions $\gamma^{E,l} = \mathcal{E}(p_1, \ldots, l, \ldots, p_n)$, the implied row degree restrictions $\boldsymbol{\gamma}^{R,l} = \mathcal{R}(p_1, \ldots, l, \ldots, p_n)$, and compare these to the existing SSVS indicators $\boldsymbol{\gamma}^S$ with respect to (7); if any discrepancies occur, we must set $\Pr\left(p_i = l \,|\, \mathbf{p}_{-i}, \boldsymbol{\gamma}_i^S, \mathbf{f}, \Lambda_{ii}, \mathbf{y}_i\right) = 0$. The correct conditional distribution then becomes

$$\Pr\left(p_i = l \,|\, \mathbf{p}_{-i}, \boldsymbol{\gamma}^S, \mathbf{f}, \Lambda_{ii}, \mathbf{y}_i\right) \propto \begin{cases} 0 \text{ if } \gamma_{j,ik}^{B,E,l} = 0, \gamma_{j,ik}^{B,R,l} \neq 0, \gamma_{j,ik}^{B,S} \neq 0 \text{ for any } i,j,k \\ p(\mathbf{y}_i \,|\, p_i = l, \boldsymbol{\gamma}_i^S, \mathbf{f}, \Lambda_{ii}) \text{ otherwise} \end{cases}.$$

(8)

Since this results in the echelon form being enforced at every iteration, draws of the Kronecker indices are equivalent to the draws of the row degrees, i.e. $\boldsymbol{\kappa} = \mathbf{p}$.

This step underscores the way in which the algorithm relies on SSVS shrinkage of all coefficients to move across possible echelon forms. To further highlight the intuition, consider once more the VARMA$(1, 1)$ example of subsection 2.2. Suppose that at a given Gibbs iteration, the model is in echelon form with $p_1 = 1, p_2 = 1$. From Table 3, it is evident that the current draw of $\boldsymbol{\gamma}^S$ must have $\gamma_{0,21}^{B,S} = 0$. Suppose it is also the case that $\gamma_{1,21}^{B,S} = 1$ (the values of the remaining indicators are irrelevant for this example) and consider sampling p_1 conditional on p_2 and the SSVS indicators. If we propose a change to $p_1 = 0$, the resulting system would assume an echelon form characterized by $p_1 = 0, p = 1$, which according to Table 3 necessitates $\gamma_{1,21}^{B,S} = 0$. Conditional on the fact that the current value $\gamma_{1,21}^{B,S} = 1$, therefore, implies that $\Pr\left(p_1 = 0 \,|\, p_2, \boldsymbol{\gamma}^S, \mathbf{f}, \Lambda_{ii}, \mathbf{y}_i\right) = 0$, and so the system must remain with $p_1 = 1$.

Suppose at the next iteration of Step 2, we sample a new $\gamma_{1,21}^{B,S}$ conditional on $p_1 = 1, p_2 = 1$. Given these row degrees, the prior for $\gamma_{1,21}^{B,S}$ is unrestricted and as long as

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$\Pr(\gamma_{1,21}^{B,S} = 1) \neq 1$, there is a positive probability of obtaining a draw of $\gamma_{1,21}^{B,S} = 0$. Once this occurs, two implications immediately follow. First, the resulting system is a sparse representation of a VARMA$_E(1,1)$ (i.e. with more zero restrictions than strictly required for identification). Second, it opens the sampler to the possibility of switching to a different echelon form, namely VARMA$_E(0,1)$.

Indeed, when we return to the sampling of p_1 conditional on p_2 and $\gamma_{1,21}^{B,S} = 0$, we now obtain a positive $\Pr\left(p_1 = 0 \,|\, p_2, \boldsymbol{\gamma}^S, \mathbf{f}, \Lambda_{ii}, \mathbf{y}_i\right) > 0$. However, if we would specify $\Pr(\gamma_{1,21}^{B,S} = 1) = 1$, then the sampler would never be able to switch to a VARMA$_E(0,1)$, regardless of how the chain is initialized. If we further set $\Pr(\gamma_{1,12}^{B,S} = 1) = 1$, then the only possible echelon forms are VARMA$_E(0,0)$ and VARMA$_E(1,1)$, but the sampler would never be able to switch between the two. In this sense, sparse echelon form structures provide the bridges by which the sampler is able to traverse the various echelon forms.

Another important feature of (8) is that it still only requires the evaluation of partial likelihoods $p(\mathbf{y}_i | \;\cdot\;)$. At first glance, this may seem counter-intuitive since a change p_i generates changes in restrictions across different equations. The reasoning is the following. Conditional on $\mathbf{f}$, each equation is independent and the likelihood can be factored as

$$p\left(\mathbf{y} \,|\, \mathbf{p}, \boldsymbol{\gamma}^S, \mathbf{f}, \boldsymbol{\Lambda}\right) = p\left(\mathbf{y}_i \,|\, p_i, \boldsymbol{\gamma}_i^S, \mathbf{f}, \Lambda_{ii}\right) \prod_{k \neq i} p\left(\mathbf{y}_k \,|\, p_k, \boldsymbol{\gamma}_k^S, \mathbf{f}, \Lambda_{kk}\right).$$

Now, in sampling p_i conditional on $\mathbf{p}_{-i}$, any proposed value of $p_i = l$ would only generate a different value of $p\left(\mathbf{y}_k \,|\, p_k, \boldsymbol{\gamma}_k^S, \mathbf{f}, \Lambda_{kk}\right)$, for $k \neq i$, if it implies a change in some SSVS indicator pertaining to equation k, i.e. $\gamma_{j,km}^{B,S}$. However, as discussed above, this would automatically imply that the conditional distribution $\Pr\left(p_i = l \,|\, \cdot\right) = 0$ (and hence, there is no need to evaluate the likelihood).

Alternatively, for any proposed value of $p_i = l$ that does not generate a different $\gamma_{j,km}^{B,S}$ for any $k \neq i$, $m = 1, \ldots, n$, and $j = 1, \ldots, \kappa_{\max}$, the quantity $\prod_{k \neq i} p\left(\mathbf{y}_k \,|\, \cdot\right)$ is unaffected, and therefore, it is constant for all possible values of p_i with nonzero weights. In consequence, it will drop out in the normalization of the weights, and hence, need not be computed for sampling purposes. Indeed, the fact that only conditional likelihoods need to be computed for sampling $p_1, \ldots, p_n$ represents another important computational advantage of this Gibbs algorithm, particularly for large dimensional systems.[11]

Implementing Step 2 is straightforward, although care must be taken in two regards: (i) respecting the restrictions implied by echelon form (when the echelon algorithm is used) on the SSVS indicators and (ii) sampling the SSVS indicators and coefficients in the correct order. The way in which we proceed depends on whether hard SSVS or soft SSVS priors are in effect. For hard SSVS priors, we sample each indicator $\gamma_{j,ik}^{\cdot,S}$ conditional

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[11]We have tried a number of alternative specifications in which $\boldsymbol{\gamma}^E = \mathcal{E}(p_1, \ldots, p_n)$ are used to impose the echelon form and the SSVS indicators $\boldsymbol{\gamma}^S$ are independent of $p_1, \ldots, p_n$. We consistently found the resulting MCMC algorithms to be computationally inferior to the one presented above, particularly in large dimensions. For example, drawing $p_1, \ldots, p_n$ jointly using an M-H proposal leads to prohibitively high rejection rates in models with $n > 3$; sampling p_i conditional on $\mathbf{p}_{-i}$ requires evaluating the joint collapsed likelihood $p(\mathbf{y} | \;\cdot\;)$ up to $n\kappa_{\max}$ times for each Gibbs run, and likewise leads to an insurmountable computational burden in large models. The specification we use (that relies on row degrees and SSVS indicators to enforce the echelon form through (7)) appears to strike an optimal balance between computational burden and mixing efficiency.

on all other indicators in equation i, but marginally of $\mathbf{B}_{(i)}, \mathbf{\Phi}_{(i)}$ (again benefiting from that fact that $\mathbf{B}, \mathbf{\Phi}$ can be integrated out analytically conditional on $\mathbf{f}$ and the indicators). Observe that this leads to a partially collapsed Gibbs sampler (van Dyk and Park, 2008), where all row degrees and SSVS indicators are blocked together with the coefficients, but we use further conditioning steps to sample the row degrees and SSVS indicators marginal of the coefficients.

In this case, it is important that the SSVS indicators are sampled prior to the coefficients $\mathbf{B}_{(i)}, \mathbf{\Phi}_{(i)}$ in order to preserve the correct target distribution. Furthermore, if the echelon form is being imposed, then conditional on $p_1, \ldots, p_n$ we must respect (7) and leave at zero any $\gamma_{j,ik}^{B,S}$ corresponding to $\gamma_{j,ik}^{B,E} = 0$ and $\gamma_{j,ik}^{B,R} = 1$. Otherwise, proceed by computing weights for a Bernoulli draw of $\gamma_{j,ik}^{B,S}$ (and similarly for $\gamma_{j,ik}^{\Phi,S}$) that are proportional to $p(\mathbf{y}_i \mid \cdot)$. The collapsed likelihood is evaluated in a fashion similar to Step 1, with specific details provided in the online appendix.

When the soft SSVS priors are specified, the conventional approach is to sample $\gamma_{j,ik}^{B,S}$ conditional on $B_{j,ik}$ and $\gamma_{j,ik}^{\Phi,S}$ conditional on $\Phi_{j,ik}$. In this case, there is no need to evaluate the likelihood, but the SSVS indicators are no longer sampled marginal of the coefficients; instead, conditional on the coefficients, the SSVS indicators are independent of each other. An important implication is that this leads to a different type of partially collapsed Gibbs sampler, where only the row degrees are blocked with the coefficients $\mathbf{B}, \mathbf{\Phi}$. Therefore, with soft SSVS priors it is crucial to sample the coefficients $\mathbf{B}, \mathbf{\Phi}$ before the SSVS indicators. Otherwise, sampling $\gamma_{j,ik}^{B,S}$ conditional on $B_{j,ik}$ (and likewise $\gamma_{j,ik}^{\Phi,S}$ conditional on $\Phi_{j,ik}$) is straightforward (again, see the online appendix for details). For the exact echelon algorithm, constraints dictated by (7) must once more be respected and $\gamma_{j,ik}^{B,S}$ maintained at zero where required.

We conclude by noting that the Gibbs algorithms described above can be used for selecting identifying restrictions or deciding whether individual coefficients are zero or not. Of course, alternative methods of model comparison, involving marginal likelihoods or information criteria can be done using MCMC output. However, in high dimensional multivariate time series models involving latent variables (such as our SSVS prior involves) marginal likelihoods can be difficult to calculate using MCMC methods, being unstable unless huge numbers of MCMC draws are used (see Chan and Grant, 2015). Marginal likelihoods are also often sensitive to the prior. In our empirical section, we use predictive likelihoods and the Deviance Information Criterion (DIC) for model comparison (see Chan and Grant, 2014). The online appendix includes more details about the DIC, including definitions and explanations of how we calculate it.

3.4 Extensions

In our empirical work, we use the models described in the preceding sub-sections. However, we note that many extensions are possible. In this sub-section, we describe two directions which may be of use for the empirical macroeconomist. The first is to allow for a time-varying $\mathbf{\Omega}_t$ or $\mathbf{\Sigma}_t$. This can be done in a standard way by adding appropriate blocks to the MCMC algorithm. For instance, multivariate stochastic volatility of the form used in Primiceri (2005) can be included by adding the extra blocks to the MCMC

algorithm as described in Appendix A of his paper.

A second extension we consider is related to an alternative approach to analyzing medium and large datasets. Specifically, let $\mathbf{y}_t$ be an $n \times 1$ vector of dependent variables that is categorized as follows:

- $\mathbf{y}_{1,t}$: the n_1 variables of primary interest;

- $\mathbf{y}_{2,t}$: the n_2 variables that together with $\mathbf{y}_t$ constitute a full $n_1 + n_2$ variate VARMA process;

- $\mathbf{y}_{3,t}$: the n_3 additional variables that are used to identify factors $\mathbf{f}_t$.

Then, consider the following expanded form representation of the VARMA model:

$$\mathbf{y}_t = \sum_{j=1}^{p} \mathbf{B}_j \mathbf{y}_{t-j} + \sum_{j=0}^{q} \mathbf{\Phi}_j \mathbf{f}_{t-j} + \boldsymbol{\eta}_t, \qquad \mathbf{f}_t \sim \mathcal{N}(\mathbf{0}, \mathbf{\Omega}_t) \text{ and } \boldsymbol{\eta}_t \sim \mathcal{N}(\mathbf{0}, \mathbf{\Lambda}), \qquad (9)$$

where $\mathbf{\Phi}_0, \ldots, \mathbf{\Phi}_q$ are $n \times n_1$ coefficient matrices and $\mathbf{f}_t$ is $n_1 \times 1$. Consequently, the covariance matrix $\mathbf{\Lambda}$ is of dimension $n \times n$, whereas the time-varying covariance matrix $\mathbf{\Omega}_t$ is diagonal with diagonal elements $\exp(h_{1,t}), \ldots, \exp(h_{n_1,t})$, where the log-volatilities follow a random walk process.

When $n \gg n_1$, (9) becomes a dynamic factor model, or under certain restrictions, a factor-augmented vector autoregression (FAVAR). In this case, identification is achieved without the need for echelon form restrictions. For example, consider the following dynamic factor representation:

$$\begin{aligned} \mathbf{y}_t &= \mathbf{\Xi}(L)\mathbf{f}_t + \boldsymbol{\zeta}_t, & \mathbf{f}_t &\sim \mathcal{N}(\mathbf{0}, \mathbf{\Omega}_t), \\ \mathbf{B}(L)\boldsymbol{\zeta}_t &= \boldsymbol{\epsilon}_t, & \boldsymbol{\epsilon}_t &\sim \mathcal{N}(\mathbf{0}, \mathbf{\Lambda}), \end{aligned}$$

where $\Xi(L) = B(L)^{-1}\Phi(L)$. Since by construction $\Xi_0 = \Phi_0$ is lower triangular with ones on the diagonal and Ω_t is diagonal, the rotation of factors and loadings is fixed and this dynamic factor model is fully identified (e.g., Bai and Wang, 2012). If $B(L)$ and $\Xi(L)$ are identified then so is $\Phi(L)$, or equivalently, pre-multiplying by $B(L)$ to obtain (9) preserves identification, thus eliminating the need for further restrictions. However, the SSVS prior for the $B(L)$ and $\Phi(L)$ coefficients can be maintained to ensure parsimony.

4   Empirical Results

In the online appendix, we investigate the performance of our algorithms using artificial data sets of relatively small sample size ($T = 100$) and VARMAs of varying dimensions up to 12. We found our algorithms generally to work well. However, MCMC efficiency deteriorated somewhat for our echelon algorithm for large dimensional models particularly when hard SSVS was used. For this reason, we argue that our approximate row degree algorithm may be useful in large models. However, when using soft SSVS, MCMC mixing was still fairly good even in VARMAs with 12 dependent variables. Hence, in this section

we will use our echelon form algorithm with soft SSVS. In the online appendix, we compare the echelon form and row degree algorithms in our largest VARMA using the macroeconomic data set described below. On the whole, we find that they are yielding similar results, but we present evidence that the row degree algorithm is failing to pick up roughly 20% of the restrictions implied by the echelon form.

In this section, we investigate the performance of our echelon algorithm (using soft SSVS and the prior specified in the online appendix) in a substantive empirical application involving quarterly US macroeconomic data in VARMAs of varying dimensions: $n = 3$, $n = 7$ and $n = 12$. In all cases we set $\kappa_{\max} = 4$ (i.e. $p = q = 4$). The justification for this, rather large, choice for lag length is that we hope our algorithm can act as a lag length selection mechanism. That is, we hope that the researcher using our approach can routinely choose fairly large values for p and q and trust the algorithm to select a parsimonious correct lag structure (which may vary across equations). In the first sub-section, we present results for this preferred model. In the following sub-section, we compare our preferred model to a conventional VARMA and a VAR. To be precise, our conventional VARMA$(4, 4)$ is the semi-structural VARMA in (1) with $\mathbf{B}_0 = \mathbf{\Theta}_0 = \mathrm{I}$, using SSVS shrinkage priors on the VAR and MA coefficients, but without considering any echelon form restrictions. Our VAR has four lags and SSVS priors on the VAR coefficients. It is obtained by starting with the echelon form VARMA, discarding the echelon restrictions, and setting $q = 0$.

Our data covers the quarters 1959:Q1 to 2013:Q4. As is commonly done (e.g., Stock and Watson, 2008) and recommended in Carriero, Clark and Marcellino (2011), each series is transformed to stationarity. We use a recursive identification scheme for our impulse responses following standard practice when working with large macroeconomic data sets (e.g. Bernanke, Boivin, and Eliasz, 2005, and Banbura, Giannone and Reichlin, 2010). In particular, we treat the Federal Funds rate as the monetary policy instrument (which is orthogonal to all other shocks) and classify every other variable as either "slow-moving" or "fast-moving" relative to this. Variables are ordered as slow-moving, then the monetary policy instrument, then the fast-moving variables. We stress that our variables have been transformed (e.g. interest rates and GDP is log differenced) and that impulse responses reported below are to these transformed variables. Exact definitions of the variables, their transformations and classifications are given in the Data Appendix.

4.0.1 Results for our Preferred Model

In this sub-section, we focus on our preferred approach described above. We run the algorithm for 50,000 iterations (5,000 burn-in) for the $n = 3$ model, 200,000 iterations (20,000 burn-in) for the $n = 7$ model, and 1,000,000 iterations (100,000 burn-in) for the $n = 12$ model.[12] For each model, we then thin the chains to obtain 10,000 draws (i.e.., for $n = 3$ we take every 5th draw, for $n = 7$ every 20th draw and for $n = 12$ every 100th draw).

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[12]Computation time on an i5-3210M dual-core, 2.50Ghz, 4GM RAM computer takes roughly 90 minutes per 100,000 draws for the $n = 12$ VARMA. For the $n = 7$ and $n = 3$ VARMAs comparable times are 45 minutes and 20 minutes. However, as the dimension of the model increases, the number of draws required to achieve a desired level of accuracy increases due to the slower mixing of the MCMC algorithm.
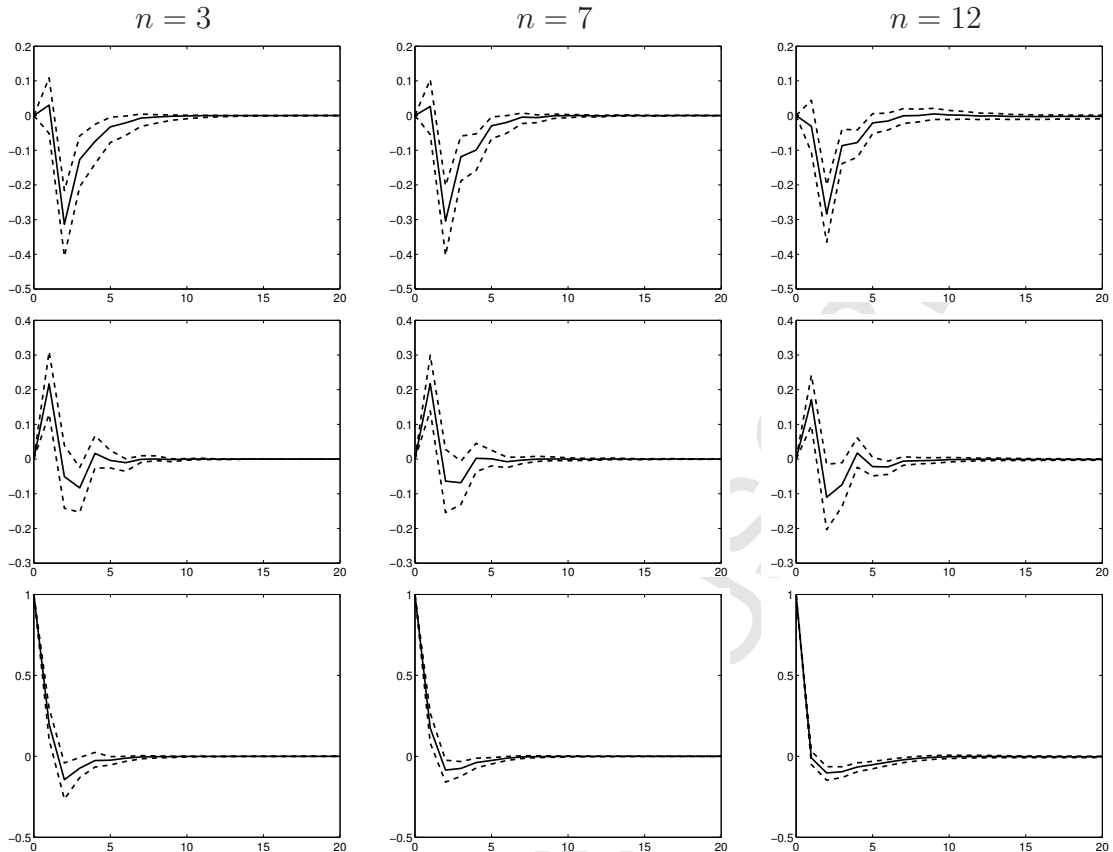
Figure 1: Impulse responses to a shock in the interest rate. The first row contains responses of GDP to a shock in the interest rate; the second row contains responses of inflation to a shock in the interest rate; the third row contains responses the interest rate to its own shock. The dotted lines depict the $(10\%, 90\%)$ HPD intervals.

Figure 1 presents the estimated impulses responses of GDP, inflation and the interest rate to a shock in the interest rate, for 20 quarters following the shock. Table 4 presents inefficiency factors relating to these impulse responses. Specifically, it contains summary statistics for the inefficiency factors of the 60 different impulse responses computed. These summary statistics indicate that the number of draws taken is longer than necessary if one is only interested in obtaining impulse responses.

Since we are interested in accurately estimating the Kronecker indices, we also present results on MCMC performance relating to them. However, since $\kappa_1, \dots, \kappa_n$ are discrete random variables, inefficiency factors are not an appropriate way to gauge sampling efficiency. In addition, any particular κ_i may naturally exhibit little movement over the course of the sampler. For instance, if there is one correct choice for κ_i then a good MCMC sampler would often (or even always) make such a choice and a lack of switching in the chain could be consistent with good MCMC performance. Accordingly, we shed light on the efficiency of the algorithm by the number of times the sampler switches

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Table 4: Comparison of inefficiency factors for impulse responses across the three models: $n = 3$, $n = 7$, and $n = 12$; note that the reported inefficiency factors are computed on thinned draws.

n	IF avg	IF st dev	IF max
3	5.90	3.17	16.10
7	1.86	2.38	15.22
12	1.17	0.43	2.90

models, as defined by the entire vector $\boldsymbol{\kappa}$. Specifically, we compute the metric

$$\varpi_n = \sum_{g=1}^{G} \mathbb{1}\left(\sum_{i=1}^{n} \left|\kappa_i^{(g)} - \kappa_i^{(g-1)}\right| > 0\right)/G,$$

where G is the number of MCMC draws, and consider that 10% represents sufficient mobility for estimation purposes. This metric is reported in Table 5 along with the estimated $\boldsymbol{\kappa}$ for the VARMAs of different dimensions. Two general points are worth noting: the MCMC sampler is mixing well and the identification restrictions selected are much more parsimonious than the VARMA(4,4) estimating model. These facts suggest our modelling approach and associated MCMC algorithm are working well, even in large VARMAs.

In the online appendix, we also present evidence on the precise values for κ visited by the MCMC sampler for the 12-variate VARMA. Six values for κ receive more than one percent of the posterior probability, but no single value receives more than fifty percent. The posterior mode of κ is similar to the posterior mean presented in Table 5.

It is worth noting that, for output and inflation, the estimated Kronecker indices are consistent across VARMAs of different dimensions. In contrast, the Kronecker index for the interest rate decreases as the size of the system increases. This result is related to the ordering of the variables and is, in fact, consistent with the Kronecker index theory. Loosely speaking, a Kronecker index κ_i represents a threshold beyond which autocovariances of further lags are linearly dependent on the lower-degree autocovariances of variables $1, \ldots, i$. Since output and inflation are always ordered first, we expect that the associated Kronecker indices do not change as additional variables are introduced. However, moving from three variables to seven, and especially from seven to twelve, introduces new variables that precede the interest rate. The fact that the Kronecker index on the interest rate shrinks from an estimated $\hat{\kappa}_3 = 1.68$ for the $n = 3$ system to $\hat{\kappa}_9 = 0.01$ for the $n = 12$ system indicates that the additional variables contain all necessary information to explain the autocorrelations present in the interest rate. In other words, we infer from the $n = 12$ system that the interest rate only responds contemporaneously to slow moving variables; removing these variables from the model leads us to estimate the interest rate as an autocorrelated process.

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Table 5: Comparison of estimated Kronecker indices across the three models: $n = 3$, $n = 7$, and $n = 12$

		n = 3	n = 7	n = 12
1	Real Gross Domestic Product	2.05	1.99	2.00
2	Consumer Price Index: All Items	2.12	2.01	2.00
3	Real Personal Consumption Exp.			1.00
4	Housing Starts: Total			1.00
5	Average Hourly Earnings: Manuf.		3.00	3.00
6	Real Gross Private Domestic Invest.			1.00
7	All Employees: Total nonfarm			1.00
8	ISM Manuf.: PMI Composite Index			1.00
9	Effective Federal Funds Rate	1.68	0.99	0.01
10	S&P 500 Stock Price Index		1.00	0.84
11	M2 Money Stock		1.28	0.97
12	Spot Oil Price: West Texas Interm		0.88	0.40
ϖ_n		23.8%	13.7%	13.4%

The preceding table suggests our methodology is successfully picking out parsimonious identified models. This issue can be investigated more deeply by looking at the estimates of the VARMA coefficients. In the online appendix, we present these for the $n = 12$ model. For comparison, this appendix also presents the estimates of autoregressive coefficients obtained from the 12-variate VAR(4). The estimates of AR and MA coefficients are mostly zeros, particularly at longer lag lengths. This strengthens the evidence that our algorithm is successfully achieving parsimony. However, we find several non-zero coefficients in $\mathbf{\Theta}_1$ (and some in $\mathbf{\Theta}_2$) indicating that adding MA terms to the VAR is important. A careful examination of the MA coefficients shows that it is usually errors in the housing starts and the purchasing manager's index equations that are found to be important. It is interesting to note that these two variables are typically regarded as leading indicators. Results for the housing starts variable are particularly interesting. When estimating the VARMA, we are finding in most equations that housing starts' effect is best modelled through the MA part of the model. That is, other variables typically react to innovations in the housing starts equation, not lags of the housing starts variable itself (i.e. the VAR coefficients relating to the housing starts variables are mostly zeros). Of course, the VAR itself could not produce such a finding. It is interesting to note that in the VAR lagged housing starts now appear much more prominently, included in some equations at the second or third lag. This is as theory would predict. A parsimonious VARMA, such as that we are finding, may be approximated by a VAR. However, the resulting VAR will be less parsimonious and with a longer lag length.

4.0.2 Comparison with Alternative Approaches

In order to investigate the advantages of working with a VARMA over a VAR and the importance of imposing identification, in this sub-section we compare our preferred approach to a different VARMA (which does have prior shrinkage but does not explicitly

Table 6: Estimated DIC values and associated numerical standard errors (in parentheses).

	$n = 3$	$n = 7$	$n = 12$
VARMA$_E(\boldsymbol{\kappa})$	1654.8	3738.1	4674.3
	(0.46)	(0.56)	(0.64)
VARMA(4,4)	1645.5	3748.1	4685.5
	(0.38)	(0.16)	(0.27)
VAR(4)	1654.5	3763.8	4738.9
	(0.12)	(0.08)	(0.10)

Table 7: Sum of log predictive likelihoods for VARMA$_E(4, 4)$, VARMA(4,4) and VAR(4).

	$n = 3$	$n = 7$	$n = 12$
VARMA$_E(4, 4)$	-182.5	-401.9	-492.3
VARMA(4,4)	-188.1	-406.0	-504.2
VAR(4)	-187.1	-406.7	-496.9

impose identification) and a VAR (which does have shrinkage but no MA components). In particular, for each model of dimension n, we compare the following specifications:

- VARMA$_E(\boldsymbol{\kappa})$: our preferred echelon form VARMA with soft SSVS priors on AR and MA coefficients and $\kappa_{\max} = 4$;

- VARMA$(4, 4)$: a VARMA with soft SSVS priors but no echelon form restrictions;

- VAR(4): a VAR with soft SSVS priors.

Note that we are only comparing modelling approaches which involve prior shrinkage. As we shall discuss below, empirical results such as impulse responses are clearly inferior and imprecise when we do not do have such shrinkage.

We begin by calculating DICs and predictive likelihoods for each model and report the results in Tables 6 and 7. The predictive likelihoods are based on the last ten years of data. It can be seen that our VARMA$_E(\boldsymbol{\kappa})$ is the preferred model by a substantial margin for models of all dimensions. The one exception to this is when $n = 3$, DIC is indicating that the VARMA$(4, 4)$ is preferred.

Each column in Tables 6 and 7 contains results for a different value of n and, thus, a different y_t. Hence, results are not comparable across columns and the tables cannot be used to provide evidence for or against working with a large dimensional model. In order to discuss the relative merits of models of different dimension, Table 8 presents predictive likelihoods for the variables which are common to all models. This allows for a comparison of different dimensional VARMAs and VARs, at least in terms their ability to forecast inflation, output growth and the interest rate. The large VARMA$_E(\boldsymbol{\kappa})$ with $n = 12$ is forecasting best of all the models and dimensions. With either VARMA approach we are finding the worst forecast performance for the $n = 3$ model, with larger dimensional models having higher predictive likelihoods. This finding does not hold for

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Table 8: Sum of log predictive likelihoods for $\text{VARMA}_E(4,4)$, VARMA(4,4) and VAR(4) based on the predictive density of the three variables in the $n = 3$ case.

	$n = 3$	$n = 7$	$n = 12$
$\text{VARMA}_E(4,4)$	-182.5	-182.2	-181.1
VARMA(4,4)	-188.1	-185.4	-187.4
VAR(4)	-187.1	-187.2	-191.0

the VAR where forecast performance deteriorates when we move away from the smallest VAR.

Next we compare impulse responses for the different models and choices for n.

Figures 2, 3 and 4 plot conventional impulse responses of our three main variables to a monetary policy shock. Our findings of the preceding sub-section indicate that the housing starts variable is found to be of particular importance and, in this section, we have evidence in favor of $n = 12$. Accordingly, in Figure 5, we plot impulse responses relating to this variable for different models for $n = 12$. The overall message of these figures, and Figure 5 in particular, is that MA components and identification can have an appreciable impact on impulse responses.

If we compare $\text{VARMA}_E(\boldsymbol{\kappa})$, VARMA(4,4) and VAR(4) impulse responses in Figures 2, 3 and 4, we see some differences in the point estimates. The impulse responses produced by the $\text{VARMA}_E(\boldsymbol{\kappa})$ are slightly smoother, having less of the irregular up and down movements of the impulse responses produced by the other approaches, particularly for $n = 12$. Furthermore, the HPD intervals are tighter when using the $\text{VARMA}_E(\boldsymbol{\kappa})$.

However, more substantive differences between the three approaches are found in Figure 5. This figure plots impulse responses relating to the housing variable for the $\text{VARMA}_E(\boldsymbol{\kappa})$, VARMA(4,4) and VAR(4). We can see the benefits of the $\text{VARMA}_E(\boldsymbol{\kappa})$ in that it is producing smooth and sensible point estimates of impulse responses with fairly tight HPD intervals about them. The VARMA(4,4) and VAR(4) are producing slightly more irregular impulse responses and the HPD intervals are wider. These differences could lead to different policy conclusions. Looking at the results generated by the $\text{VARMA}_E(\boldsymbol{\kappa})$ model, it appears that the interest rate will continue to increase[13] in response to a housing starts shock, even after 20 quarters. This finding is significant in the sense that the HPD interval is entirely above zero. The housing start variable itself is very slow in adjusting downward following a positive shock, suggesting that increasing interest rates exerts little effect in discouraging further real-estate expansion. This is partly confirmed by the impulse response of housing starts to an increase in the interest rate. That is, after 20 quarters the model predicts a negative impact with a high degree of certainty (e.g. the HPD interval is all below zero), but one that is very small in magnitude—i.e., approximately -0.025 on average after 20 quarters.

We do not get quite the same picture by looking at the responses generated with the VARMA(4,4) and VAR(4). This is mainly due to the larger degree of imprecision of the impulse responses of the models. For instance, with the VAR the impulse response of the interest rate to a shock in housing starts is approximately zero after 20 quarters, with

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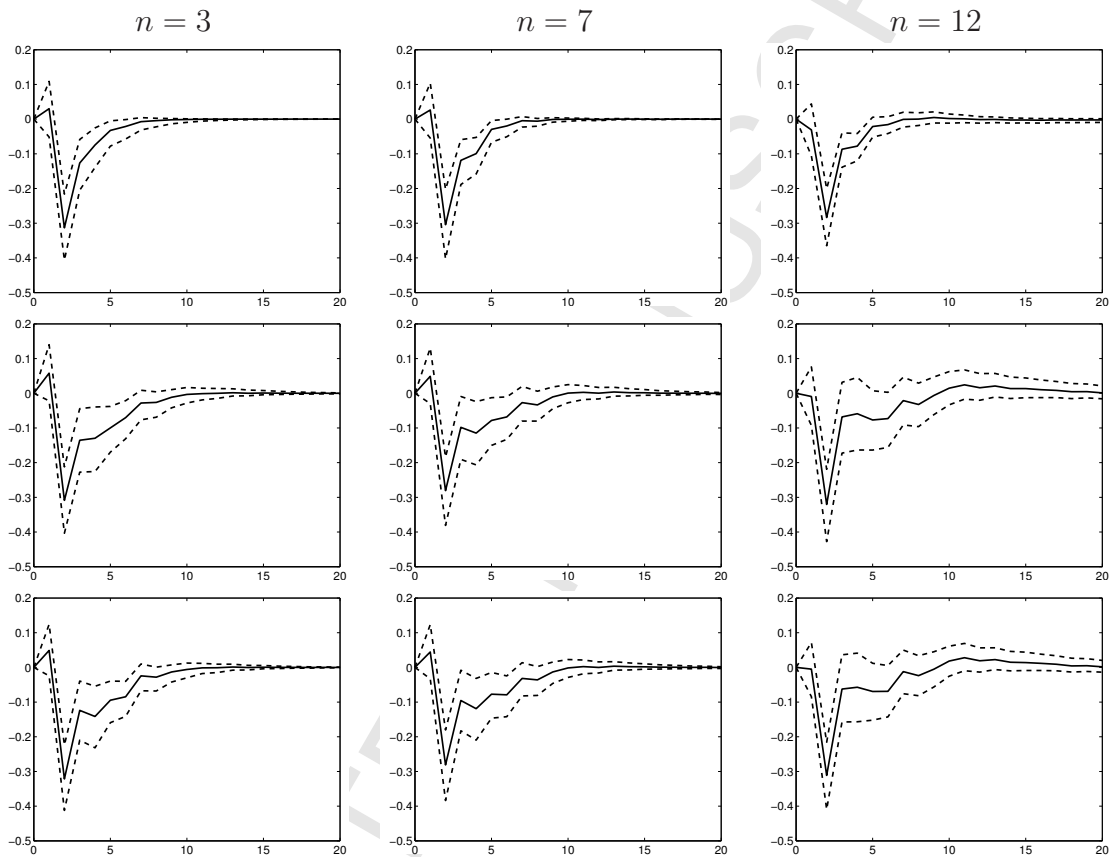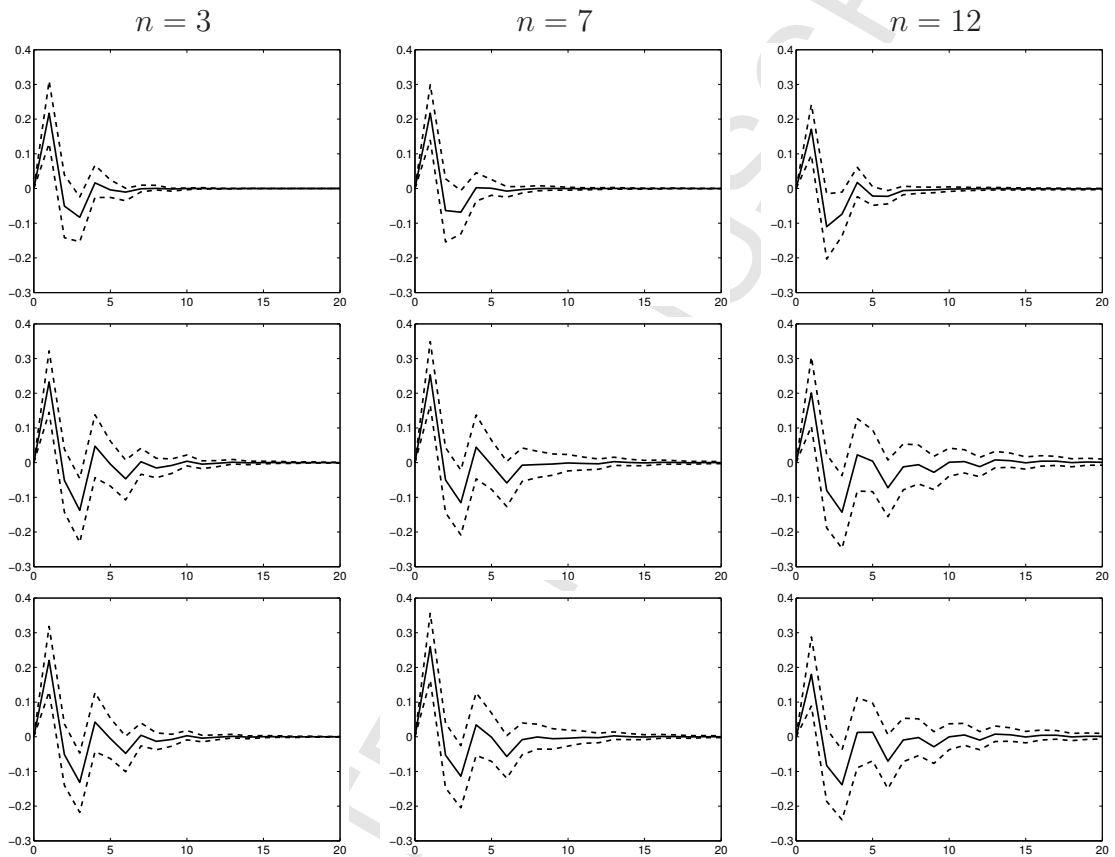[13]Recall that the interest rate series is first-differenced.

Figure 2: Comparison of impulse responses of GDP to a shock in the interest rate. The first, second and third rows contain results for the $\text{VARMA}_E(\boldsymbol{\kappa})$, $\text{VARMA}(4, 4)$ and $\text{VAR}(4)$, respectively. The dotted lines depict the $(10\%, 90\%)$ HPD intervals.

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Figure 3: Comparison of impulse responses of inflation to a shock in the interest rate. The first, second and third rows contain results for the $\text{VARMA}_E(\boldsymbol{\kappa})$, $\text{VARMA}(4,4)$ and $\text{VAR}(4)$, respectively. The dotted lines depict the $(10\%, 90\%)$ HPD intervals.
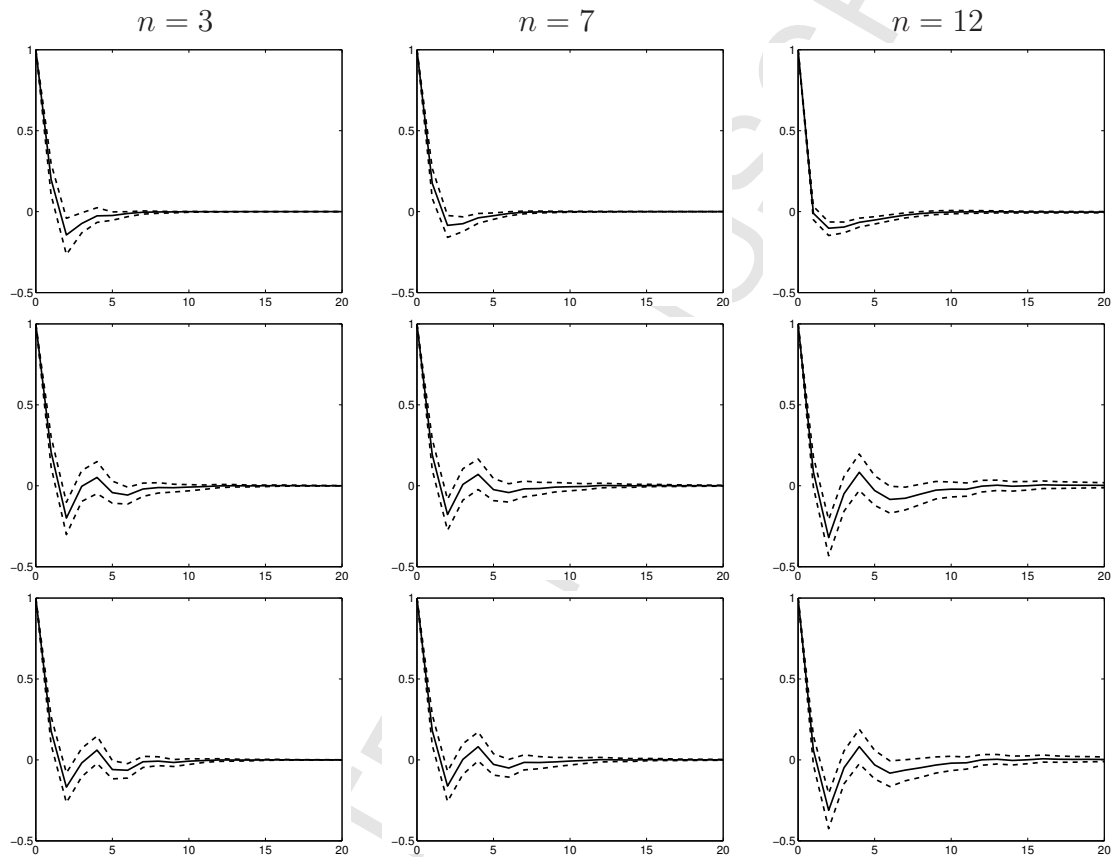
29

Figure 4: Comparison of impulse responses of the interest rate to own shock. The first, second and third rows contain results for the VARMA$_E(\boldsymbol{\kappa})$, VARMA(4, 4) and VAR(4), respectively. The dotted lines depict the $(10\%, 90\%)$ HPD intervals.
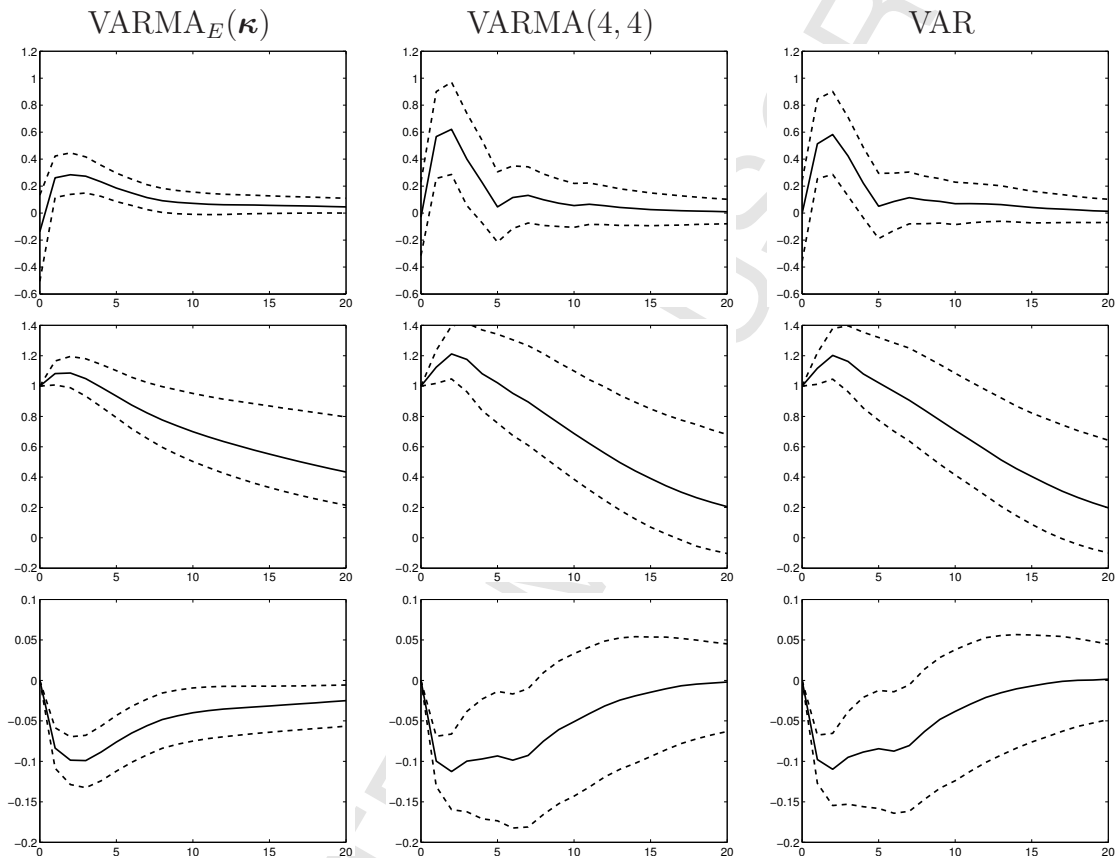
30

Figure 5: Comparison of impulse responses of the housing start and interest rate to shocks. The first row contains responses of the interest rate to a shock in the housing start; the second row contains responses of the housing start to its own shock; the third row contains responses of the housing start to a shock in the interest rate. The dotted lines depict the $(10\%, 90\%)$ HPD intervals.

the HPD interval covering both positive and negative regions. Also, the impulse response of housing starts to its own shock is initially large under the VAR, but then falls faster than what the VARMA$_E(\boldsymbol{\kappa})$ predicts. At the same time, the VAR generates responses of housing starts to an increase in the interest rate such that the median response vanishes by the end of the 20 quarter horizon. This indicates that an increase in the interest rate has no long term effect on the housing starts, although the HPD intervals are substantially wider than those produced by the VARMA.

All of the approaches discussed so far in this sub-section have included shrinkage using SSVS priors. If we do not include such shrinkage, impulse responses become even more irregular and HPD intervals become even wider. For the sake of brevity, we will not produce conventional impulse responses similar to Figures 2 through 4 for VARMAs and VARs without shrinkage. Suffice it to note here that there is an appreciable deterioration in impulse responses relative to Figures 2 through 4. Instead Figure 6 presents impulse responses without prior shrinkage relating to the housing variable for $n = 12$. Relative to the VARMA$_E(\boldsymbol{\kappa})$ both the VARMA$(4,4)$ and VAR(4) are producing impulse responses which are much more erratic and with much wider HPD intervals.
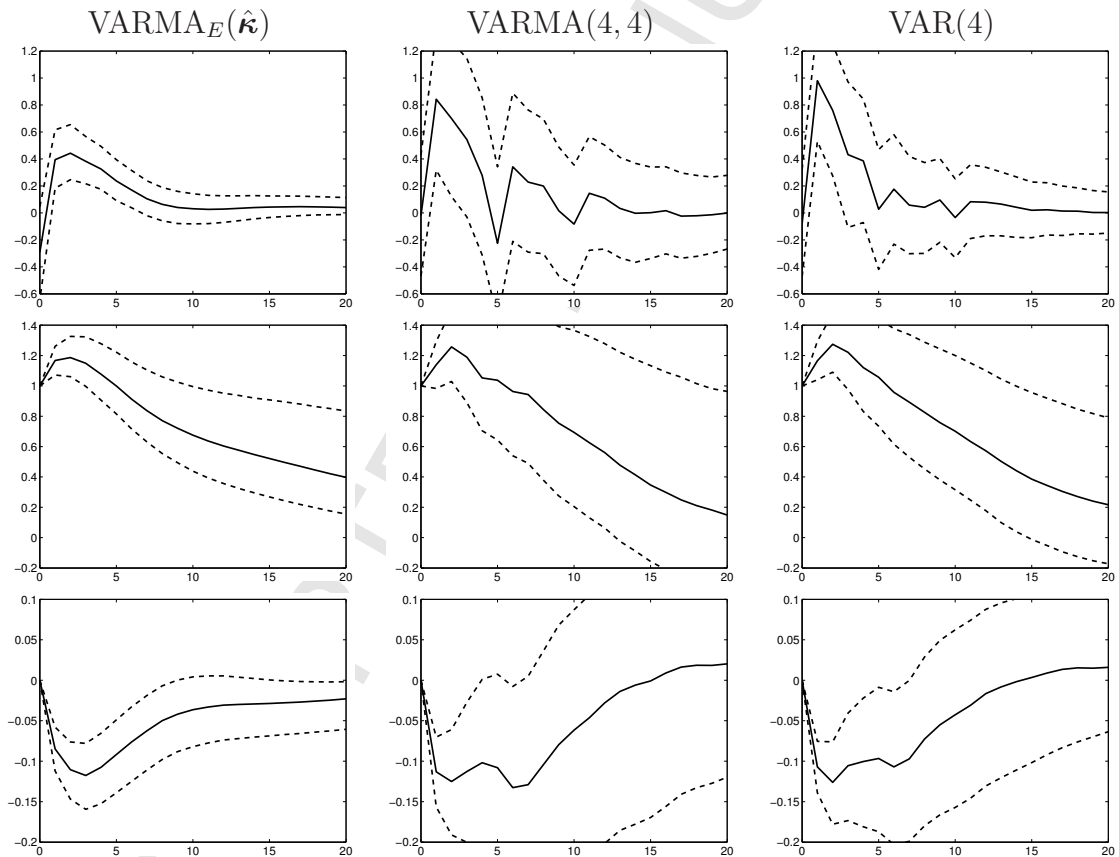

Figure 6: Comparison of impulse responses of the housing start and interest rate to shocks, without using SSVS shrinkage. The first row contains responses of the interest rate to a shock in the housing start; the second row contains responses of the housing start to its own shock; the third row contains responses of the housing start to a shock in the interest rate. The dotted lines depict the $(10\%, 90\%)$ HPD intervals.

In sum, the specification, identification and shrinkage issues investigated in this paper can have an important impact on policy-relevant issues.

5 Conclusions

We began this paper by arguing that there might be some benefits to working with VARMAs instead of VARs. However, VARMAs are little-used due to problems of identification, over-parameterization and computation. In this paper, we have developed a modelling approach, using SSVS priors on both parameters and identification restrictions, which surmounts these problems. In a substantive macroeconomic application, we show that this modelling approach does work well even in VARMAs of high dimension. It is computationally feasible and yields sensible results which have the potential to lead to different policy conclusions than simpler VAR or alternative VARMA approaches.

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Data Appendix

All variables were downloaded from St. Louis' FRED database and cover the quarters 1959:Q1 to 2013:Q4. The following table lists the variables, describes how they were transformed and whether they are slow- or fast-moving variables. The transformation codes are: 1 - no transformation (levels); 2 - first difference, 3 - second difference; 4 - logarithm; 5 - first difference of logarithm; 6 - second difference of logarithm.

Variable	Trans. Code	Slow / Fast	included in model $n = 3$	$n = 7$	$n = 12$
Real Gross Domestic Product	5	S	X	X	X
Consumer Price Index: All Items	6	S	X	X	X
Real Personal Consumption Exp.	5	S			X
Housing Starts: Total	4	S			X
Average Hourly Earnings: Manuf.	6	S		X	X
Real Gross Private Domestic Invest.	5	S			X
All Employees: Total nonfarm	5	S			X
ISM Manuf.: PMI Composite Index	1	S			X
Effective Federal Funds Rate	2		X	X	X
S&P 500 Stock Price Index	5	F		X	X
M2 Money Stock	6	F		X	X
Spot Oil Price: West Texas Interm.	5	F		X	X