Large Bayesian Matrix Autoregressions

Joshua C. C. Chan        Yaling Qi
Purdue University        Purdue University

Preliminary Draft
September 2023

Abstract

High-dimensional matrix-valued time-series are increasingly common in economics and finance. Prominent examples include large cross-region panels and dynamic economic networks. As the dimensions of the matrix grow, conventional approaches based on vector autoregressions—implemented by vectoring the matrix-valued data—become computationally infeasible. We introduce a class of large Bayesian matrix autoregressions (MARs) that can accommodate time-varying volatility, non-Gaussian errors and COVID-19 outliers. To tackle parameter proliferation, we propose Minnesota-type shrinkage priors for these MARs. We develop a unified approach for estimating these models that scales well to high dimensions. The empirical relevance of these new MARs is illustrated using a US state-level dataset that contains 6 macroeconomic times-series for each of the 50 states, with a total of 300 times-series.

JEL classification: C11, C32, C55

Keywords: heavy-tailed distribution, outlier, shrinkage prior, stochastic volatility, tensor, vector autoregression
1 Introduction

Matrix-valued data observed over time are common in economics, finance and related areas. A classic example is a cross-country panel dataset in which a few key macroeconomic indicators for each country are observed over time (Canova and Ciccarelli, 2009, 2013; Koop and Korobilis, 2016). More recently, larger cross-region panels with more regional units and economic variables, such as state-level or other sub-national level time-series datasets, have become widely available (Baumeister, Leiva-León, and Sims, 2022; Bokun, Jackson, Kliesen, and Owyang, 2023; Koop, McIntyre, Mitchell, Poon, and Wu, 2023). Another fast growing category of large matrix-valued time-series are dynamic economic networks, such as bilateral trade volumes among trading partners (Kharrazi, Rovenskaya, and Fath, 2017; Kapetanios, Serlenga, and Shin, 2021) and bilateral outstanding credits between countries (Billio, Casarin, Iacopini, and Kaufmann, 2023).

The growing availability of these complex datasets presents new opportunities, but it also exposes the limitations of conventional multivariate time-series econometric models. More specifically, a standard approach is to treat the matrix-valued observations over time as time-series vectors, which can then be conveniently modeled using vector autoregressions (VARs). There are, however, two disadvantages of this approach. First, vectoring the matrix-valued observation mixes its columns and rows, and consequently it disregards the topological structure of the data. Second, despite recent advances in modeling large VARs (Bańbura, Giannone, and Reichlin, 2010; Carriero, Clark, and Marcellino, 2019), it remains extremely time-consuming or practically infeasible to estimate VARs obtained by vectoring the high-dimensional matrix-valued data.

To tackle these challenges, we take up the matrix autoregression (MAR) introduced in Hoff (2015) and Chen, Xiao, and Yang (2021), which regresses the matrix-valued observation on its lagged values using a bilinear form. The MAR has an equivalent representation as a parsimonious VAR, where the matrix structure of the data is exploited to construct the VAR coefficient matrices using far fewer free parameters relative to an unrestricted VAR. As such, the MAR modeling framework ameliorates the two drawbacks of the VAR approach.

We further extend the MAR framework along two directions. First, instead of assuming a time-invariant, Gaussian error distribution, we introduce a class of MARs that can ac-
commodate time-varying volatility, non-Gaussian errors and COVID-19 outliers. This is motivated by the increasing recognition of the need to allow for time-varying volatility in modeling most macroeconomic datasets (see, e.g., Cogley and Sargent, 2005; Primiceri, 2005; Sims and Zha, 2006). In fact, there is now a large body of empirical evidence that demonstrates the importance of time-varying volatility for model-fit and forecasting in small VARs (Clark, 2011; D’Agostino, Gambetti, and Giannone, 2013; Clark and Ravazzolo, 2015; Chan and Eisenstat, 2018) as well as in large VARs (Koop and Korobilis, 2013; Carriero, Clark, and Marcellino, 2016, 2019; Chan, 2023a). In addition, the unexpected extreme movements in many macroeconomic variables at the onset of the COVID-19 pandemic underline the need to allow for non-Gaussian errors and potential outliers. Clark and Mertens (2023) provide a recent review on the benefits of incorporating stochastic volatility in a wide range of applications using Bayesian VARs.

Our second contribution is to introduce Bayesian shrinkage priors and efficient estimation methods that can handle large datasets. While earlier works have focused on matrix-valued time-series of moderate sizes,¹ we are interested in high-dimensional settings in which the matrix dimensions are large. For example, in our empirical application we analyze a US state-level dataset that contains 6 macroeconomic times-series for each of the 50 US states, with a total of 300 times-series. Even though a MAR has far fewer parameters compared to an unrestricted VAR, it may still have more parameters than observations over time when the dimensions of the matrix are large. Therefore, we introduce Bayesian shrinkage priors on the MAR coefficients. These new priors are inspired by the Minnesota prior of Doan, Litterman, and Sims (1984) and Litterman (1986), and can be viewed as a generalization of the Minnesota prior to the MAR setting. There priors are conjugate and hence facilitate fast estimation. Additionally, we follow Giannone, Lenza, and Primiceri (2015) and estimate the prior hyperparameters that control the overall shrinkage strength from the data, instead of fixing them at some subjective values.

Building upon the fast sampling methods in Carriero, Clark, and Marcellino (2016) and Chan (2020), we develop a unified approach for estimation—by exploiting a certain Kronecker product structure of the likelihood implied by this family of MARs—that can

¹For example, Chen, Xiao, and Yang (2021) consider an application with a panel of 5 countries, and each has 4 economic indicators. The model is fitted using the iterative least squares and maximum likelihood estimators. Celani and Pagnottoni (2023) provide a Bayesian treatment of the MAR and consider a panel of 9 countries, each with 6 economic indicators.
drastically speed up the computations. In particular, for the matrix-valued observation $Y_t$ of size $n \times k$, sampling the MAR coefficients using conventional methods would involve $O(n^6)$ and $O(k^6)$ elementary operations. The proposed sampling approach instead can be done in computational complexity of the order $O(n^3)$ and $O(k^3)$. This orders-of-magnitude speed-up makes the proposed estimation approach suitable for fitting large datasets.

The empirical relevance of these new models is illustrated using a US state-level dataset that includes 300 times-series. Even with such a large dataset, the proposed MARs, together with the Minnesota-type shrinkage priors, can be estimated relatively quickly. The estimation results demonstrate the strong interactions between the variables across states, highlighting the importance of modeling all the state variables jointly. In addition, it is also clear that there is a spike in volatility at the onset of the COVID-19 pandemic—the error standard deviation in 2020Q2 is estimated to be between 5-7 times larger than that of regular periods. These results thus underscore the importance of allowing time-varying volatility and heavy-tailed error distributions.\(^2\)

In addition to contributing to the development of more flexible matrix autoregressions, this paper is also related to two other strands of literature. First, it contributes to the emerging literature on modeling multidimensional arrays or tensors (Leng and Tang, 2012; Lock, 2018), particularly third-order tensors. Most of the existing literature does not explicitly model the dynamics, even when one of the tensor dimensions is time. Two notable exceptions are Hoff (2015) and Billio, Casarin, Iacopini, and Kaufmann (2023); the former introduces a multilinear tensor autoregression based on the Tucker product, whereas the latter develops a more general linear autoregressive tensor process where the tensor coefficients are parameterized using a PARAFAC decomposition. In contrast to the time-invariant Gaussian error distribution considered in both papers, here we develop a framework that can accommodate non-Gaussian errors and richer time-varying dynamics.

This paper also contributes to the literature on modeling and forecasting regional data. Hamilton and Owyang (2012) is a classic paper that uses US state-level payroll employment data to infer regional recessions. Koop, McIntyre, and Mitchell (2020) and Koop,\(^2\)

---

\(^2\)A few recent papers, such as Schorfheide and Song (2021) and Lenza and Primiceri (2022) using US data and Bobeica and Hartwig (2023) using euro area data, have shown that impulse response functions and forecasts from homoskedastic VARs are heavily distorted by the extreme observations related to the COVID-19 pandemic.
McIntyre, Mitchell, and Poon (2020) develop a mixed-frequency framework to nowcast UK regional growths using both regional and national data. More recent papers such as Baumeister, Leiva-León, and Sims (2022) and Bokun, Jackson, Kliesen, and Owyang (2023) have used larger US state-level datasets for nowcasting or monitoring state-level economic conditions. Our paper provides a convenient modeling framework that can handle datasets with a large number of regional units and economic indicators.

The rest of this paper is organized as follows. Section 2 first introduces a general framework for modeling matrix-valued time-series with a flexible error covariance structure. It then offers a few different interpretations of the matrix autoregression and discusses some identification issues. Lastly, the section develops Bayesian shrinkage priors that generalize the Minnesota priors to MAR settings. Section 3 proposes a unified approach to estimate these flexible MARs using Markov chain Monte Carlo (MCMC) methods. Section 4 considers an application that involves a US state-level dataset with 300 time-series. Lastly, Section 5 concludes and outlines some future research directions.

## 2 A Flexible Framework for Matrix Autoregressions

We introduce a general framework for Bayesian matrix autoregressions that aims to strike the right balance between flexibility and tractability in high-dimensional settings. On the one hand, this flexible framework can accommodate a wide variety of empirically relevant features, including heavy-tailed error distributions, time-varying volatility and robustness to outliers. On the other hand, it also facilitates fast computation and can be used to model large datasets.

### 2.1 The Modeling Framework

To set the stage, let $\mathbf{Y}_t$ denote an $n \times k$ matrix of endogenous variables at time $t$ for $t = 1, \ldots, T$. To fix ideas, one may think of each column of $\mathbf{Y}_t$ containing the $n$ variables for each of the $k$ regions. In our empirical application that models US state-level data, we have $k = 50$ states and each state has $n = 6$ variables, with a total of 300 variables.

A common approach to model the matrix-valued data $\mathbf{Y}_t$ is to first stack its columns into
a vector, i.e., vec\(\left(Y_t\right)\), which is then fitted using the vector autoregression with \(p\) lags:

\[
vec\left(Y_t\right) = \Phi_1 vec\left(Y_{t-1}\right) + \cdots + \Phi_p vec\left(Y_{t-p}\right) + e_t, \tag{1}
\]

where \(\Phi_1, \ldots, \Phi_p\) are \(nk \times nk\) coefficient matrices and \(e_t\) is an \(nk \times 1\) vector of errors. There are two main drawbacks of modeling \(Y_t\) using the VAR in (1). First, by vectoring \(Y_t\), the columns and rows of \(Y_t\) are mixed. Consequently, the VAR ignores the matrix structure—e.g., the strong connections between the variables in the same region (column) and those between the same variable (row) across regions. The second drawback is the proliferation of parameters when either \(n\) or \(k\) is large. For example, for \(n = 6, k = 50\) and \(p = 2\), there are 180,000 VAR coefficients, which makes estimation and inference practically infeasible.

To tackle these two issues, we follow Hoff (2015) and Chen, Xiao, and Yang (2021) to directly model the evolution of the matrix \(Y_t\) via the following matrix autoregression (MAR):

\[
Y_t = A_1 Y_{t-1} B_1' + \cdots + A_p Y_{t-p} B_p' + E_t, \tag{2}
\]

where \(A_1, \ldots, A_p\) and \(B_1, \ldots, B_p\) are, respectively, \(n \times n\) and \(k \times k\) coefficient matrices. For simplicity we exclude the intercepts; a matrix of intercepts or any deterministic term can be easily added to the model. The above bilinear form facilitates model interpretation and estimation. In particular, the matrix autoregression in (2) can be represented in the form of a VAR:

\[
vec\left(Y_t\right) = (B_1 \otimes A_1) vec\left(Y_{t-1}\right) + \cdots + (B_p \otimes A_p) vec\left(Y_{t-p}\right) + vec\left(E_t\right),
\]

where \(\otimes\) denotes the Kronecker product. Hence, the MAR can be viewed as a special case of the VAR, where the VAR coefficient matrix is modeled as the Kronecker product \(\Phi_j = (B_j \otimes A_j)\). Consequently, the number of VAR coefficients is reduced from \(n^2k^2p\) to \((n^2 + k^2)p\). Subsection 2.2 provides more discussions on the interpretation of the MAR and its relations to the VAR.

While earlier works consider only homoskedastic MARs where the distribution of the \(n \times k\) matrix of errors, \(E_t\), is time-invariant, we propose a more general setting in which
$E_t$ has a conditionally Gaussian distribution given the latent variable $\omega_t$:

$$\text{vec}(E_t) \sim \mathcal{N}(0_{nk}, \omega_t \Sigma_c \otimes \Sigma_r), \quad (3)$$

where $\Sigma_c$ and $\Sigma_r$ are, respectively, $k \times k$ and $n \times n$ covariance matrices. The homoskedastic MAR considered in Hoff (2015) and Chen, Xiao, and Yang (2021) can be recovered as a special case with $\omega_1 = \cdots = \omega_T = 1$. By assuming different distributions for the mixing variables $\omega_1, \ldots, \omega_T$, this framework encompasses a wide range of flexible error distributions that are found empirically useful for modeling macroeconomic and financial data. Below we give a few important examples.

1. **Heavy-tailed distributions.** Since many distributions can be represented as a scale mixture of normals, the conditionally Gaussian specification in (3) can accommodate many common heavy-tailed distributions that are useful to capture rare but large changes in volatility. For example, if the mixing variable $\omega_t$ follows the inverse-gamma distribution $\omega_t \sim \mathcal{IG}(\nu/2, \nu/2)$, then the marginal distribution of $\text{vec}(E_t)$ unconditional on $\omega_t$ has a multivariate $t$ distribution with zero mean, scale matrix $\Sigma_c \otimes \Sigma_r$ and degree of freedom parameter $\nu$. Alternatively, if $\omega_t$ has a gamma distribution, then marginally $\text{vec}(E_t)$ has a multivariate normal-gamma distribution, which includes the multivariate Laplace distribution as a special case. Both of these distributions have heavier tails than normals, and they generally provide better fit for data with infrequent volatility jumps. Empirical studies that find heavy-tailed errors useful in the context of VARs include Clark and Ravazzolo (2015), Cross and Poon (2016) and Chiu, Mumtaz, and Pinter (2017).

2. **Robustness to outliers.** The conditionally Gaussian specification in (3) can also be used for addressing potential outliers using a tailored mixing distribution. An important example is an explicit outlier component of the type proposed in Stock and Watson (2016). More specifically, let $\omega_t = o_t^2$ and $o_t$ follows a mixture of two distributions: a point mass at 1 and a uniform distribution on the interval $(2, 20)$. The former can be thought of as ‘regular’ observations with scale normalized to 1, whereas the latter captures ‘outliers’ that have 2-20 times larger standard deviations than regular observations. As demonstrated in Carriero, Clark, Marcellino, and Mertens (2022), this outlier component is especially useful for modeling observations associated with the COVID-19 pandemic.

3. **Time-varying volatility.** One of the most robust empirical findings in modeling macroeconomic data is the importance of allowing for time-varying volatility (e.g., Sims...
and Zha, 2006; Clark, 2011; Chan and Eisenstat, 2018). The conditionally Gaussian framework in (3) can accommodate certain types of time-varying volatility processes. An important example is the common stochastic volatility model introduced in Carriero, Clark, and Marcellino (2016). In particular, let \( \omega_t = e^{h_t} \), and assume that the log-volatility \( h_t \) follows a stationary AR(1) process with 0 mean:

\[
h_t = \phi h_{t-1} + u_t^h, \quad u_t^h \sim \mathcal{N}(0, \sigma_h^2),
\]

for \( t = 2, \ldots, T \), where \(|\phi| < 1\) and the initial condition is specified as \( h_1 \sim \mathcal{N}(0, \sigma_h^2/(1 - \phi^2)) \). The log-volatility \( h_t \) here may be interpreted as the level of economy-wide macroeconomic uncertainty (see also Jurado, Ludvigson, and Ng, 2015). Another example is the volatility model with a deterministic break date considered in Lenza and Primiceri (2022), which is designed to model the drastic increase in volatility at the onset of the COVID-19 pandemic and the subsequent gradual decrease in volatility. Their model can also be parameterized using the conditionally Gaussian framework.

Naturally, any combinations of the above heavy-tailed errors and volatility processes can also be incorporated using the conditionally Gaussian framework. For instance, one may consider a MAR with the common stochastic volatility and the outlier component. In that case, \( \omega_t = e^{h_t} o_t^2 \), where \( h_t \) follows the AR(1) process in (4) and \( o_t \) follows the two-component mixture described above. Other models such as those in Chan (2020) and Hartwig (2021) can also be considered.

While the modeling framework in (2)–(3) is flexible and includes many empirically useful specifications as special cases, it is crucial to recognize its limitations. In particular, the latent variable \( \omega_t \) is assumed to scale the entire covariance matrix of \( E_t \), implying that each element of \( E_t \) is impacted equally by \( \omega_t \). As such, the proposed framework does not nest, for example, a model in which each row of \( Y_t \) has its specific stochastic volatility factor. Estimation of such a model, however, would be practically infeasible in high-dimensional settings. The proposed framework therefore provides the right balance between modeling flexibility and computational tractability.
2.2 Model Interpretation and Identification

In this section we discuss the interpretation of the coefficient matrices in the MAR and some identification issues. For ease of exposition, we consider the case with only one lag:

\[ Y_t = A_1 Y_{t-1} B_1' + E_t. \]

In the above bilinear form, the coefficient matrix \( A_1 \) corresponds to row-wise relationships, whereas \( B_1 \) represents column-wise interactions. To tease out the impact of the two matrices, it is useful to consider a few special cases. Recall that the MAR may be viewed as a special VAR in which the VAR coefficient matrix is parameterized as \( \Phi_1 = (B_1 \otimes A_1) \). If we assume \( B_1 = I_k \), then we can express the MAR as:

\[ \text{vec}(Y_t) = (I_k \otimes A_1) \text{vec}(Y_{t-1}) + \text{vec}(E_t). \]

In other words, each column of \( Y_t \) follows the same VAR with the coefficient matrix \( A_1 \), and there are no interactions among the columns (in the conditional mean). Similarly, for the special case with \( A_1 = I_n \), each row of \( Y_t \) follows a VAR with the same coefficient matrix.

The covariance matrix of \( \text{vec}(E_t) \) has a similar interpretation. For simplicity, set \( \omega_t = 1 \). Then, the matrix of errors as specified in (3) can be equivalently represented as \( E_t = \Sigma^2_r Z_t \Sigma^2_c \), where \( Z_t \) is an \( n \times k \) matrix consisting of independent standard normal random variables. It is clear from this representation that \( \Sigma_r \) corresponds to row-wise covariances and \( \Sigma_c \) represents column-wise covariances. In particular, if \( \Sigma_c = I_k \), then \( E_t = \Sigma^2_r Z_t \), which implies that the columns of \( E_t \) are all mutually independent and each row has the same covariance matrix \( \Sigma_r \). More generally, the covariance between the \((i_1, j_1)\) and \((i_2, j_2)\) elements of \( E_t \) is \( \text{cov}(e_{t,i_1,j_1}, e_{t,i_2,j_2}) = \sigma_{r,i_1,i_2} \sigma_{c,j_1,j_2} \).

Another interpretation of the MAR is related to the global VAR (Pesaran, Schuermann, and Weiner, 2004) and the multivariate autoregressive index model (Carriero, Kapetanios, and Marcellino, 2016). More specifically, let \( y_{t,i,j}, a_{1,i,j} \) and \( b_{1,i,j} \) denote the \((i, j)\) elements of \( Y_t, A_1 \) and \( B_1 \), respectively. Then, \( y_{t,i,j} \) can be expressed as

\[
y_{t,i,j} = \sum_{l_1=1}^{n} \sum_{l_2=1}^{k} a_{1,i,l_1} b_{1,j,l_2} y_{t-1,l_1,l_2} + e_{t,i,j} = \sum_{l_1=1}^{n} a_{1,i,l_1} z_{t-1,l_1,j} + e_{t,i,j},
\]
where $z_{t-1,l_1,j} = \sum_{l_2=1}^{k} b_{1,j,l_2} y_{t-1,l_1,l_2}$ is a linear combination of the $l_1$-th row of $Y_{t-1}$ across the columns. Under this representation, one can view the MAR as a multi-equation regression with covariates constructed from linear combinations of the columns of $Y_{t-1}$. In particular, using our running state-level data example, the MAR can be interpreted as first constructing linear combinations of GDP, unemployment, etc., across states, and use them as regressors.

Next, we discuss some identification issues that arise in the MAR. First, the parameters $A_1$ and $B_1$ are not separately identified, but they are identified up to scale. That is, if $(B_1 \otimes A_1)z = (\tilde{B}_1 \otimes \tilde{A}_1)z$, for all $z$, then $\tilde{A}_1 = c A_1$ and $\tilde{B}_1 = c^{-1} B_1$ for some $c \neq 0$, provided that neither $A_1$ and $B_1$ is the zero matrix. Hence, to fix the scale, we normalize the $(1,1)$ element of $B_1$ to be 1. More generally, for the MAR of order $p$, we set the $(1,1)$ element of $B_j$ to be 1, i.e., $b_{j,1,1} = 1, j = 1, \ldots, p$. Similarly, the covariances $\Sigma_r$ and $\Sigma_c$ are only identified up to scale. We normalize the $(1,1)$ element of $\Sigma_c$ to be 1: $\sigma_{c,1,1} = 1$.

### 2.3 Bayesian Shrinkage Priors

We are interested in settings when $n$ or $k$ (or both) is large. In those cases, the matrix autoregression has a large number of parameters, and consequently regularization or shrinkage is vital for obtaining sensible results. In addition, to facilitate fast estimation, we extend the natural conjugate prior (see, e.g., Koop and Korobilis, 2010; Karlsson, 2013) designed for VARs to our setting. To that end, let $A = (A_1, \ldots, A_p)'$ and $B = (B_1, \ldots, B_p)'$, so that $A$ and $B$ are of dimensions $np \times n$ and $kp \times k$, respectively. We consider the prior of the form $p(A, B, \Sigma_r, \Sigma_c | \kappa_A, \kappa_B) = p(A, \Sigma_r | \kappa_A)p(\Sigma_c | \kappa_B)$, where $\kappa_A$ and $\kappa_B$ are some hyperparameters which we treat as unknown.

First, we assume that $(A, \Sigma_r)$ has a normal-inverse-Wishart distribution (see, e.g., Kadiyala and Karlsson, 1997; Koop and Korobilis, 2010):

$$
\Sigma_r \sim \mathcal{IW}(\nu_r, S_r), \quad (\text{vec}(A) | \Sigma_r, \kappa_A) \sim \mathcal{N}(\text{vec}(A_0), \Sigma_r \otimes V_A),
$$

where $\text{vec}(A_0)$ is the prior mean vector and the $np \times np$ prior covariance matrix $V_A$ is assumed to be diagonal and depend on the unknown hyperparameter $\kappa_A$. The joint
density function of \((A, \Sigma_r)\) is thus given by
\[
p(A, \Sigma_r) \propto |V_A|^{-\frac{n}{2}}|\Sigma_r|^{-\frac{\nu_r + n + np + 1}{2}} e^{-\frac{1}{2}tr(\Sigma^{-1}S_r)} e^{-\frac{1}{2}tr(\Sigma^{-1}(A-A_0)V^{-1}_A(A-A_0))},
\]
where \(tr(\cdot)\) is the trace operator.

We calibrate \(A_0\) and \(V_A\) in the spirit of the Minnesota priors pioneered by Doan, Litterman, and Sims (1984) and Litterman (1986). More specifically, \(\text{vec}(A_0)\) is set to be a zero vector for growth rates data. This reflects the prior belief that growth rates data are typically not very persistent, and the coefficient matrix \(A\) is thus shrunk to 0. For levels data, \(\text{vec}(A_0)\) is set to be zero except for the coefficients associated with the first own lag, which are set to be one. This expresses the preference for a random walk specification, reflecting the prior belief that levels data are generally highly persistent.

To calibrate the diagonal elements of \(V_A\), let \(\hat{s}^2_{i,*} = \sum_{j=1}^k \hat{s}^2_{i,j} / k\), where \(\hat{s}^2_{i_1,i_2}\) denotes the sample variance of an AR(4) model for the variable \(y_{t,i_1,i_2}\), the \((i_1, i_2)\) element of \(Y_t\). Hence, \(\hat{s}^2_{i,*}\) is the average sample variances of the variables in the \(i\)-th row. Then, the \(j\)-th diagonal element of \(V_A\) is assumed to be \(v_{A,j,j} = \kappa_A/((l^2\hat{s}^2_{i,*})\) for a coefficient associated with lag \(l\) of the variable in the \(i\)-th row. Intuitively, the prior variance is scaled by \(\hat{s}^2_{i,*}\), and the coefficient associated to a lag \(l\) variable is shrunk more heavily to zero as the lag length increases. The overall shrinkage strength is controlled by the hyperparameter \(\kappa_A\), where a smaller value indicates more aggressive shrinkage. We follow the recommendation of Giannone, Lenza, and Primiceri (2015) to estimate \(\kappa_A\) from the data instead of fixing it at some commonly-used subjective value. Finally, we set \(\nu_r = n + 2\), \(S_r = \text{diag}(\hat{s}^2_{1,*}, \ldots, \hat{s}^2_{n,*})\).

These hyperparameters are elicited in the spirit of the Minnesota priors. In particular, for \(k = 1\), they reduce to those of the standard Minnesota priors (see, e.g., Karlsson, 2013; Carriero, Clark, and Marcellino, 2015).

Similarly, we consider the following normal-inverse-Wishart prior on \((B, \Sigma_c)\):
\[
\Sigma_c \sim \mathcal{IW}(\nu_c, S_c), \quad (\text{vec}(B) | \Sigma_c, \kappa_B) \sim \mathcal{N}(\text{vec}(B_0), \Sigma_c \otimes V_B),
\]
where \(\text{vec}(B_0)\) is the prior mean vector and the \(kp \times kp\) prior covariance matrix \(V_B\) is assumed to be diagonal and depend on the unknown hyperparameter \(\kappa_B\). Naturally, one can elicit \(B_0\) and \(V_B\) to incorporate prior beliefs specific to the application. Below we provide a baseline case that is expected to be applicable to a wide range of cross-region
applications.

As discussed in Section 2.2, when \( B_j = I_k, j = 1, \ldots, p \), then each column of \( Y_t \), representing observations from a particular region, follows the same VAR with the coefficient matrix \( A \), and there are no interactions among the columns in the conditional mean. We therefore set the prior mean to be \( B_0 = (I_k, \ldots, I_k)' \), and shrink \( B \) toward this simpler setting. This choice of prior mean is also consistent with the identification restrictions that the \((1,1)\) elements of \( B_1, \ldots, B_p \) are 1. To calibrate the diagonal elements of \( V_B \), let \( \hat{s}_{ij}^2 = \frac{\sum_{i=1}^n s_{ij}^2}{n} \) denote the average sample variances of the variables in the \( j \)-th column. Then, the \( i \)-th diagonal element of \( V_B \) is assumed to be \( v_{B_{ij}} = \kappa_B/\left(l^2\hat{s}_{ij}^2\right) \) for a coefficient associated with lag \( l \) of the variable in the \( j \)-th column. Hence, a coefficient is shrunk more strongly to zero if it corresponds to a variable of higher lag, and the prior variance is scaled by \( \hat{s}_{ij}^2 \). The hyperparameter \( \kappa_B \) determines the overall shrinkage strength, which is again estimated from the data. We set \( \nu_c = k + 2 \), \( S_c = \text{diag}(1, \hat{s}_{2,2}/\hat{s}_{1,1}, \ldots, \hat{s}_{k,k}/\hat{s}_{k,1}) \). Here we normalize the scale matrix \( S_c \) so that it is consistent with the identification restriction that the \((1,1)\) element of \( \Sigma_c \) is fixed at one.

Finally, the hyperparameters \( \kappa_A \) and \( \kappa_B \) are assumed to have hierarchical gamma priors: \( \kappa_A \sim \mathcal{G}(c_{A,1}, c_{A,2}) \) and \( \kappa_B \sim \mathcal{G}(c_{B,1}, c_{B,2}) \).

### 3 Bayesian Estimation

In this section we provide a general discussion on the estimation of the Bayesian MARs specified in (2)-(3). In particular, we develop a fast and simple approach to sample the pairs \( (A, \Sigma_r) \) and \( (B, \Sigma_c) \) given the shrinkage hyperparameters \( \kappa = (\kappa_A, \kappa_B)' \) and an arbitrary vector of latent variables \( \omega = (\omega_1, \ldots, \omega_T)' \). In Appendix A we take up various examples of \( \omega \) and provide estimation details for tackling each case, as well as the sampling steps for \( \kappa \).

We first derive the likelihood function implied by (2)-(3). Letting \( A = (A_1, \ldots, A_p)' \) and \( B = (B_1, \ldots, B_p)' \), note that one can rewrite the mean equation in (2) as:

\[
Y_t = A'X_tB + E_t,
\]
where $X_t = \text{diag}(Y_{t-1}, \ldots, Y_{t-p})$ is an $np \times kp$ block-diagonal matrix of lagged values. Given the covariance structure in (3), the likelihood function can be expressed as:

$$p(Y \mid A, B, \Sigma_c, \Sigma_r, \omega) = (2\pi)^{-\frac{Tr(\Sigma)}{2}}|\Sigma_c|^{-\frac{T}{2}}|\Sigma_r|^{-\frac{T}{2}} \prod_{t=1}^{T} \omega_t^{-\frac{nk}{2}} e^{-\frac{1}{2\omega_t} tr(\Sigma_c^{-1}(Y_t-A'X_tB)\Sigma_r^{-1}(Y_t-A'X_tB))}. $$

(6)

Assuming the natural conjugate priors for $(A, \Sigma_r)$ and $(B, \Sigma_c)$, posterior draws can be obtained by sequentially sampling from: 1) $p(A, \Sigma_r \mid Y, B, \Sigma_c, \kappa, \omega)$; 2) $p(B, \Sigma_c \mid Y, A, \Sigma_r, \kappa, \omega)$; 3) $p(\kappa \mid Y, A, B, \Sigma_r, \Sigma_c, \omega)$; and 4) $p(\omega \mid Y, A, B, \Sigma_r, \Sigma_c, \kappa)$. Depending on how one models the latent variables $\omega$, additional blocks might be needed to sample some additional hierarchical parameters. These steps are typically easy to implement as they amount to fitting a univariate time-series model. A variety of examples are given in Appendix A. Below we provide details on implementing Step 1 and Step 2 of sampling from the high-dimensional densities $p(A, \Sigma_r \mid Y, B, \Sigma_c, \kappa, \omega)$ and $p(B, \Sigma_c \mid Y, A, \Sigma_r, \kappa, \omega)$ efficiently.

More specifically, recall that $A$ is of dimensions $np \times n$, and sampling $A$ using conventional methods would involve $O(n^6)$ elementary operations. Fortunately, it can be shown that $(A, \Sigma_r \mid Y, B, \Sigma_c, \kappa, \omega)$ has a normal-inverse-Wishart distribution, and one can sample $A$ with computational complexity of the order $O(n^3)$. To see this, note that it follows from (5) and (6) that

$$p(A, \Sigma_r \mid Y, B, \Sigma_c, \kappa, \omega) \propto \Sigma_r^{-\frac{Tr+n+p+Tk+1}{2}} e^{-\frac{1}{2} tr(\Sigma_r^{-1}S_r)}$$

$$\times e^{-\frac{1}{2} tr(\Sigma_r^{-1}(A-A_0)/V_A^{-1}(A-A_0)+\sum_{t=1}^{T} \omega_t^{-1}(Y_t-A'X_tB)\Sigma_r^{-1}(Y_t-A'X_tB'))}$$

$$=|\Sigma_r|^{-\frac{Tr+n+p+Tk+1}{2}} e^{-\frac{1}{2} tr(\Sigma_r^{-1}S_r)} e^{-\frac{1}{2} tr(\Sigma_r^{-1}(A_0V_A^{-1}A_0+\sum_{t=1}^{T} \omega_t^{-1}Y_t\Sigma_c^{-1}Y_t'\tilde{A}K_A\tilde{A}'))}$$

$$\times e^{-\frac{1}{2} tr(\Sigma_r^{-1}(A-A_0)'K_A(A-A_0))},$$

where

$$K_A = V_A^{-1} + \sum_{t=1}^{T} \omega_t^{-1}X_tB\Sigma_c^{-1}B'X_t', \quad \tilde{A} = K_A^{-1} \left( V_A^{-1}A_0 + \sum_{t=1}^{T} \omega_t^{-1}X_tB\Sigma_c^{-1}Y_t \right).$$

13
In the above derivation we have used the fact that

\[(A - A_0)'V_A^{-1}(A - A_0) + \sum_{t=1}^{T} \omega_t^{-1}(Y_t - A'X_tB)'\Sigma_c^{-1}(Y_t - A'X_tB)'
\]

\[= (A - \hat{A})'K_A(A - \hat{A}) + A_0'V_A^{-1}A_0 + \sum_{t=1}^{T} \omega_t^{-1}Y_t\Sigma_c^{-1}Y_t' - \hat{A}'K_A\hat{A}.
\]

In other words, \((A, \Sigma_r | Y, B, \Sigma_c, \kappa, \omega)\) has the normal-inverse-Wishart distribution with parameters \(\nu_r + Tk, \hat{S}_r, \hat{A}\) and \(K_A^{-1}\), where

\[\hat{S}_r = S_r + A_0'V_A^{-1}A_0 + \sum_{t=1}^{T} \omega_t^{-1}Y_t\Sigma_c^{-1}Y_t' - \hat{A}'K_A\hat{A}.
\]

Hence, we can sample \((A, \Sigma_r | Y, B, \Sigma_c, \kappa, \omega)\) in two steps. First, we sample \(\Sigma_r\) marginally from \((\Sigma_r | Y, B, \Sigma_c, \kappa, \omega) \sim IW(\hat{S}_r, \nu_r + Tk)\). Then, given the \(\Sigma_r\) drawn, we sample

\[(\text{vec}(A) | Y, B, \Sigma_r, \Sigma_c, \kappa, \omega) \sim N\left(\text{vec}(\hat{A}), \Sigma_r \otimes K_A^{-1}\right).
\]

Since the covariance matrix is of dimension \(n^2p \times n^2p\), and sampling from this high-dimensional density using conventional methods would involve \(O(n^6p^3)\) operations. This can be computationally intensive when \(n\) is large. Instead, we adopt an efficient algorithm to sample from the matrix-normal distribution to our setting (e.g., Bauwens, Lubrano, and Richard, 1999, p.320). This algorithm has been used in Carriero, Clark, and Marcellino (2016) and Chan (2020) to estimate various large Bayesian VARs. More specifically, we exploit the Kronecker structure \(\Sigma_r \otimes K_A^{-1}\) to speed up computation. Consequently, the complexity of the problem can be drastically reduced to \(O(n^3p^3)\) operations. We further improve upon this approach by avoiding the computation of the inverse of the \(np \times np\) matrix \(K_A\). The computational details are provided in Appendix A.

Similarly, it can be shown that \((B, \Sigma_c | Y, A, \Sigma_r, \kappa, \omega)\) has a normal-inverse-Wishart
distribution with parameters \( \nu_c + Tn, \hat{S}_c, \hat{B} \) and \( K_B^{-1} \), where

\[
K_B = V_B^{-1} + \sum_{t=1}^{T} \omega_t^{-1} X_t' A \Sigma_r^{-1} A' X_t, \quad \hat{B} = K_B^{-1} \left( V_B^{-1} B_0 + \sum_{t=1}^{T} \omega_t^{-1} X_t' A \Sigma_r^{-1} Y_t \right),
\]

\[
\hat{S}_c = S_c + B_0' V_B^{-1} B_0 + \sum_{t=1}^{T} \omega_t^{-1} Y_t' \Sigma_r^{-1} Y_t - \hat{B}' K_B \hat{B}.
\]

Again, we can sample \((B, \Sigma_c | Y, A, \Sigma_r, \kappa, \omega)\) in two steps. First, we sample \(\Sigma_c\) marginally from \((\Sigma_c | Y, A, \Sigma_r, \kappa, \omega) \sim IW(\hat{S}_c, \nu_c + Tn)\) with the normalization restriction that \(\sigma_{c,1,1} = 1\). This can be done using the algorithm in Nobile (2000). Then, given the sampled \(\Sigma_c\), we simulate

\[
(vec(B) | Y, A, \Sigma_r, \Sigma_c, \kappa, \omega) \sim N(vec(\hat{B}), \Sigma_c \otimes K_B^{-1})
\]

with the normalization restrictions that the (1,1) elements of \(B_1, \ldots, B_p\) are all 1. Sampling from a Gaussian distribution subjected to linear restrictions can be done efficiently by using Algorithm 2.6 in Rue and Held (2005) or Algorithm 2 in Cong, Chen, and Zhou (2017). We provide the details of this sampling step in Appendix A.

### 4 Empirical Application

To illustrate the utility of the proposed models and estimation methods, we consider an application that involves a US state-level dataset. More specifically, for each of the 50 US states, we obtain 6 quarterly time-series sourced from the Bureau of Labor Statistics and the FRED database maintained by the Federal Reserve Bank of St. Louis. These 6 variables are initial unemployment insurance claims, continued unemployment insurance claims, total nonfarm employment, unemployment rate, new housing permits, and real home price index. The sample period is from 1991Q1 to 2023Q1. A detailed description of the variables and their transformations are provided in Appendix B. We represent the data at time \( t \) as an \( n \times k \) matrix \( Y_t \), where the columns refer to the \( k = 50 \) states and the rows are the \( n = 6 \) variables.
4.1 Full Sample Results

We first report various estimates of interest using the full sample, which includes the COVID-19 pandemic. As widely noted, the COVID-19 pandemic has caused extreme movements in many macroeconomic and financial time-series, and a failure to account for these outliers would result in heavily distorted parameter estimates, as demonstrated in recent papers such as Schorfheide and Song (2021), Lenza and Primiceri (2022) and Bobeica and Hartwig (2023). Therefore, we consider two Bayesian MARs that explicitly account for time-varying volatility or potential outliers.

More specifically, both models can be nested within the proposed framework and represented as the system in (2)–(3). The first Bayesian MAR incorporates the common stochastic volatility specification proposed in Carriero, Clark, and Marcellino (2016) with \( \omega_t = e^{h_t} \), and the log-volatility \( h_t \) follows the stationary AR(1) given in (4). This model is referred to as BMAR-CSV. The second model includes the outlier component introduced in Stock and Watson (2016) with \( \omega_t = o_t^2 \), where \( o_t \) follows a 2-part distribution with a point mass at 1 and a uniform distribution on the interval (2, 20). This latter model is referred to as BMAR-O. For both models we set the lag order to be \( p = 2 \).

To visualize the correlation pattern among the variables (rows), we report in Figure 1 a heatmap of the posterior means of \( A_1 \) and \( A_2 \) from the BMAR-CSV, where red entries denote positive values, blue negative and white zero. First, it is clear that, as expected, all variables are rather persistent on average. For example, the AR(1) coefficient for nonfarm payroll is estimated to be about 0.44, and the AR(1) and AR(2) coefficient estimates for unemployment rate are, respectively, 0.63 and \(-0.36\). Second, the proposed hierarchical shrinkage prior on \( A = (A_1, A_2)' \) strongly shrinks many of the off-diagonal elements to zero. In particular, the shrinkage hyperparameter \( \kappa_A \)—that controls the overall shrinkage strength on \( A \), where a smaller value indicate more aggressive shrinkage to zero—is estimated to be 0.032 (compared to the prior mean of 1). Despite the shrinkage effects, some off-diagonal elements corresponding to closely related variables remain non-zero. For example, lagged continued claims have non-negligible impacts on initial claims, and vice versa, possibly reflecting current labor market conditions.
Figure 1: A heatmap of the posterior means of $A_1$ and $A_2$ from the BMAR-CSV.

Next, Figure 2 reports a heatmap of the posterior means of $B_1$ from the BMAR-CSV, which represents the correlation structure among the states (columns). Due to space constraint, we do not report estimate of $B_2$, but the pattern is similar. Despite the fact that the proposed hierarchical shrinkage prior is designed to shrink $B_1$ to the identity matrix, the majority of the off-diagonal elements are estimated to be non-zero. This is also reflected in the estimated shrinkage hyperparameter $\kappa_B$ that controls the overall shrinkage strength on $B = (B_1, B_2)'$: its posterior mean is about 0.84 compared to the prior mean of 1, indicating that the data do not favor strong shrinkage of the off-diagonal elements to zero. Not surprisingly, many neighboring states or states with similar outputs show stronger interactions. For instance, lagged variables of Virginia and Florida most positively impact the variables of Alabama with corresponding coefficients estimated to be 0.38 and 0.33, respectively.$^3$ Overall, these results highlight the strong interactions between the states.

$^3$In our sample the variables of Alabama are stacked in the first column. And since we normalize the scale of the MAR coefficients by setting the (1,1) elements of $B_1$ and $B_2$ to be 1, the magnitude of the two states’ variables are 38% and 33% of those of own state variables.
To assess the extent of time-varying sizes of shocks, we report the posterior means of the time-varying standard derivations from the BMAR-CSV and the BMAR-O in Figure 3. Despite the two very different modeling approaches—the BMAR-CSV prescribes a persistent volatility process whereas the BMAR-O assumes serial independence of the occurrence of the outliers—the estimated time-varying standard derivations from the two models are remarkably similar. In particular, for most of the sample before the onset of the COVID-19 pandemic in 2020Q2, the standard derivations were mostly around 1 (normalized as ‘regular’ observations). In 2020Q2, the standard derivations jumped to 5.5 for the BMAR-CSV and 7 for the BMAR-O, and they stayed elevated afterward. These
results underscore the empirical relevance of explicitly modeling time-varying volatility or allowing for outliers.

Figure 3: Posterior means of the time-varying standard derivations, $e^{h_t}$ and $o_t$ respectively, from BMAR-CVS (top panel) and BMAR-O (bottom panel).

5 Concluding Remarks and Future Research

Two recent developments have motivated our paper: the increasing recognition of the need to allow for flexible time-varying features in modeling most macroeconomic datasets and the growing availability of a large number of matrix-valued time-series. In response to these recent developments, we have introduced a new class of matrix autoregressions that can accommodate time-varying volatility, non-Gaussian errors and COVID-19 outliers. We then developed an efficient, unified approach that scales well to high-dimension datasets. We illustrated the methodology using a US state-level dataset that involves 300 macroeconomic time-series.
There are multiple lines of future research that are worth pursuing. First, it would be useful to extend the MARs to a mixed-frequency framework—e.g., modeling both quarterly and monthly time-series simultaneously. This can be done, for example, by incorporating the data augmentation approach in Schorfheide and Song (2015) or Chan, Poon, and Zhu (2023) to simulate the missing monthly observations of the quarterly data. An interesting application would be one that aims to construct monthly state-level GDP estimates using both quarterly and monthly variables. Another promising direction is to develop time-varying parameter MARs. In a VAR setting, Chan (2023b) has found evidence that the VAR coefficients in some, but not all, equations are time-varying. The binary indicator approach in Chan (2023b) can be adopted to model the time-varying MAR coefficients. In addition, both the dynamic shrinkage approach of Koop and Korobilis (2018) or the dynamic shrinkage with sparsification approach of Huber, Koop, and Onorante (2019) are promising alternatives.
Appendix A: Estimation Details

In this appendix we provide estimation details of the proposed Bayesian matrix autoregressions.

A1: Sampling A and B

For sampling the coefficient matrices $A$ and $B$—of dimensions $np \times n$ and $kp \times k$, respectively—from their full conditional distributions, we make use of some standard results on the matrix normal distribution (see, e.g., Bauwens, Lubrano, and Richard, 1999, pp. 301-302). Specifically, an $r \times s$ random matrix $Z$ is said to have a matrix normal distribution $\mathcal{MN}(M, S \otimes R)$ for covariance matrices $R$ and $S$ of dimensions $r \times r$ and $s \times s$, respectively, if and only if $\text{vec}(Z) \sim \mathcal{N}(\text{vec}(M), S \otimes R)$. Naturally, a bilinear transformation of a matrix normal random matrix followed by a translation is also a matrix normal random matrix. More precisely, suppose $Z \sim \mathcal{MN}(M, S \otimes R)$ and let $V = CZD + E$. Then, $V \sim \mathcal{MN}(CMD + E, (D'SD) \otimes (CRC'))$.

Now, we can sample $\text{vec}(A) \sim \mathcal{N}(\text{vec}(\hat{A}), \Sigma_r \otimes K_A^{-1})$ as follows. Let $C_{K_A}$ and $C_{\Sigma_r}$ be the lower Cholesky factors of $K_A$ and $\Sigma_r$, respectively. We claim that if we construct

$$A = \hat{A} + C_{K_A}^{-1'}ZC_{\Sigma_r}'$$

where $Z$ is an $np \times n$ matrix of iid $\mathcal{N}(0, 1)$ random variables, then $\text{vec}(A)$ has the desired distribution. To show that, since $Z \sim \mathcal{MN}(0, I_n \otimes I_{np})$, using the previous result with $C = C_{K_A}^{-1'}, D = C_{\Sigma_r}'$ and $E = \hat{A}$, we have $A \sim \mathcal{MN}(\hat{A}, \Sigma_r \otimes K_A^{-1})$ and therefore, by definition, $\text{vec}(A) \sim \mathcal{N}(\text{vec}(\hat{A}), \Sigma_r \otimes K_A^{-1})$. Finally, we note that in the above construction, one can efficiently compute $C_{K_A}^{-1'}Z$ by solving the linear system $C_{K_A}'X = Z$ for $X$ without explicitly obtaining the inverse $C_{K_A}^{-1}$.

Next, we outline the sampling of $\text{vec}(B) \sim \mathcal{N}(\text{vec}(\hat{B}), \Sigma_c \otimes K_B^{-1})$ with the normalization restrictions that the $(1, 1)$ elements of $B_1, \ldots, B_p$ are all one. To that end, we first represent the normalization restrictions as a system of $p$ linear restrictions: $M \text{vec}(B) = b_0$, where $M = (m_{i,j})$ is a $p \times k^2p$ selection matrix with $m_{i,(i-1)k+1} = 1$ for $i = 1, \ldots, p$, and $b_0$ is a $p \times 1$ vector of ones. Then, we can apply Algorithm 2.6 in Rue and Held.
(2005) or Algorithm 2 in Cong, Chen, and Zhou (2017) to efficiently sample \( \text{vec}(\mathbf{B}) \sim \mathcal{N}(\text{vec}(\hat{\mathbf{B}}), \Sigma_c \otimes K_B^{-1}) \) so that \( \mathbf{M} \text{vec}(\mathbf{B}) = \mathbf{b}_0 \). In particular, one can first sample \( \text{vec}(\mathbf{B}_u) \) from the unconstrained conditional posterior distribution using the algorithm discussed earlier, and construct

\[
\text{vec}(\mathbf{B}) = \text{vec}(\mathbf{B}_u) + \left( \Sigma_c \otimes K_B^{-1} \right) \mathbf{M}' \left( \mathbf{M} \left( \Sigma_c \otimes K_B^{-1} \right) \mathbf{M}' \right)^{-1} (\mathbf{b}_0 - \mathbf{M} \text{vec}(\mathbf{B}_u)).
\]

Then, \( \text{vec}(\mathbf{B}) \) has the distribution \( \mathcal{N}(\text{vec}(\hat{\mathbf{B}}), \Sigma_c \otimes K_B^{-1}) \) such that \( \mathbf{M} \text{vec}(\mathbf{B}) = \mathbf{b}_0 \). We summarize the algorithm in Algorithm 1.

**Algorithm 1** Sampling \( \mathcal{N}(\text{vec}(\hat{\mathbf{B}}), \Sigma_c \otimes K_B^{-1}) \) such that \( \mathbf{M} \text{vec}(\mathbf{B}) = \mathbf{b}_0 \).

1. Sample \( \mathbf{B}_u = \hat{\mathbf{B}} + C_{K_B}^{-1/2} \mathbf{Z} C_{\Sigma_c}^t, \) where \( \mathbf{Z} \) is a \( kp \times k \) matrix of \( \mathcal{N}(0,1) \) random variables.
2. Compute \( \mathbf{C} = C_{\Sigma_c^{-1}} \otimes C_{K_B} \), where \( C_{\Sigma_c^{-1}} \) is the lower Cholesky factor of \( \Sigma_c^{-1} \).
3. Solve \( \mathbf{CC}' \mathbf{U} = \mathbf{M}' \) for \( \mathbf{U} \)
4. Solve \( \mathbf{MUV} = \mathbf{U}' \) for \( \mathbf{V} \).
5. Return \( \text{vec}(\mathbf{B}) = \text{vec}(\mathbf{B}_u) + \mathbf{V}'(\mathbf{b}_0 - \mathbf{M} \text{vec}(\mathbf{B}_u)). \)

**A2: Sampling \( \kappa_A \) and \( \kappa_B \)**

Next, we discuss the sampling steps of drawing the hyperparameters \( \kappa_A \) and \( \kappa_B \). First, note that \( \kappa_A \) only appears in two terms: its gamma prior \( \kappa_A \sim \mathcal{G}(c_{A,1}, c_{A,2}) \) and \( \mathbf{V}_A \), the prior covariance matrix of \( \mathbf{A} \), which is an \( np \times np \) diagonal matrix with the \( i \)-th diagonal element \( v_{A,i,i} = \kappa_A C_{A,i} \) for some constant \( C_{A,i} \). Then, we can express the conditional distribution of \( \kappa_A \) as

\[
p(\kappa_A | \mathbf{A}, \Sigma_r) \propto \kappa_A^{c_{A,1}-1} \frac{1}{2} e^{-c_{A,2} \kappa_A} \times |\mathbf{V}_A|^{-\frac{n}{2}} e^{-\frac{1}{2} \text{tr} \left( \Sigma_r^{-1} (\mathbf{A}-\mathbf{A}_0)' \mathbf{V}_A^{-1} (\mathbf{A}-\mathbf{A}_0) \right)}
\]

\[
\times \kappa_A^{c_{A,1} - \frac{n^2}{2} - 1} e^{-c_{A,2} \kappa_A} \mathbf{V}_A^{-\frac{1}{2}} \text{tr} \left( \mathbf{V}_A^{-1} (\mathbf{A}-\mathbf{A}_0) \Sigma_r^{-1} (\mathbf{A}-\mathbf{A}_0)' \mathbf{V}_A^{-1} (\mathbf{A}-\mathbf{A}_0) \right)
\]

\[
\times \kappa_A^{c_{A,1} - \frac{n^2}{2} - 1} e^{-\frac{1}{2} \left( 2c_{A,2} \kappa_A + \kappa_A^{-1} \sum_{i=1}^{np} Q_{A,i,C_{A,i}} \right)}.
\]
where $Q_{A,i}$ is the $i$-th diagonal element of $Q_A = (A - A_0)\Sigma^{-1}_r (A - A_0)'$. Note that this is the kernel of the generalized inverse Gaussian distribution

$$GIG \left( c_{A,1} - \frac{n^2p}{2}, 2c_{A,2}, \sum_{i=1}^{np} Q_{A,i}/C_{A,i} \right).$$

Draws from the generalized inverse Gaussian distribution can be obtained using the algorithm in Devroye (2014).

Similarly, $\kappa_B$ only appears in its gamma prior $\kappa_B \sim G(c_B, 1, c_B, 2)$ and $V_B$, which is a $kp \times kp$ diagonal matrix where the $i$-th diagonal element is $v_{B,i,i} = \kappa_B C_{B,i}$ for some constant $C_{B,i}$. It can be shown that $(\kappa_B | B, \Sigma_c)$ has the generalized inverse Gaussian distribution:

$$GIG \left( c_{B,1} - \frac{k^2p}{2}, 2c_{B,2}, \sum_{i=1}^{kp} Q_{B,i}/C_{B,i} \right),$$

where $Q_{B,i}$ is the $i$-th diagonal element of $Q_B = (B - B_0)\Sigma^{-1}_c (B - B_0)'$.

**A3: Sampling Other Parameters**

We now consider a few specific examples of $\omega$ and discuss how one can modify the posterior sampler outlined in the main text to handle each case.

**Example 1 Student’s t errors**

As discussed in Section 2 of the main text, the case of $t$ distributed errors is nested within the proposed framework: the latent variables $\omega = (\omega_1, \ldots, \omega_T)'$ are distributed independently as $(\omega_i | \nu) \sim IG(\nu/2, \nu/2)$.

Posterior draws can be obtained by sequentially sampling from: 1) $p(A, \Sigma_r | Y, B, \Sigma_c, \kappa, \omega)$; 2) $p(B, \Sigma_c | Y, A, \Sigma_r, \kappa, \omega)$; 3) $p(\kappa | Y, A, B, \Sigma_r, \Sigma_c, \omega)$; 4) $p(\omega | Y, A, B, \Sigma_r, \Sigma_c, \kappa, \nu)$; and 5) $p(\nu | Y, A, B, \Sigma_r, \Sigma_c, \kappa, \omega)$. Steps 1-2 can be implemented exactly as described in Section 3 of the main text and Step 3 as outlined in Section A2. For Step 4, let $s_i^2 = \text{tr} (\Sigma_i^{-1} E_i \Sigma_i^{-1} E_i)$, where $E_i$ can be computed given $Y_i, A$ and $B$ using (2). Then,
the conditional distribution of $\omega$ can be expressed as:

$$p(\omega \mid Y, A, B, \Sigma_r, \Sigma_c, \kappa, \nu) = \prod_{t=1}^{T} p(\omega_t \mid Y, A, B, \Sigma_r, \Sigma_c, \kappa, \nu) \propto \prod_{t=1}^{T} \omega_t^{-(\nu+1)/2} e^{-\frac{\nu}{2\omega_t}} \times \omega_t^{-\frac{nk}{2}} e^{-\frac{1}{2\omega_t s_t^2}}.$$  

That is, each $\omega_t$ is conditionally independent given the data and other parameters, and has an inverse-gamma distribution: $(\omega_t \mid Y, A, B, \Sigma_r, \Sigma_c, \kappa, \nu) \sim IG((nk+\nu)/2, (s_t^2+\nu)/2)$.

Lastly, $\nu$ can be sampled by an independence-chain Metropolis-Hastings step with the proposal distribution $\mathcal{N}(\tilde{\nu}, K_{\nu}^{-1})$, where $\tilde{\nu}$ is the mode of $\log p(\nu \mid Y, A, B, \Sigma_r, \Sigma_c, \omega)$ and $K_{\nu}$ is the negative Hessian evaluated at the mode. For implementation details of this step, see Chan and Hsiao (2014).

**Example 2 Outlier detection**

The proposed framework can also be used to incorporate the approach in Stock and Watson (2016) and Carriero, Clark, Marcellino, and Mertens (2022) to handle potential outliers. To that end, let $\omega_t = o_t^2$, where $o_t$ follows a mixture distribution that distinguishes between regular observations with $o_t = 1$ and outliers for which $o_t > 2$. More specifically, $o_t$ equals 1 with probability $1 - p_o$; $o_t$ follows a uniform distribution on $(2, 20)$ with probability $p_o$. The outlier probability $p_o$ is assumed to have a beta prior $\mathcal{B}(a_0, b_0)$, where the hyperparameters $a_0$ and $b_0$ are calibrated so that the mean outlier frequency is once every 4 years in quarterly data.

Posterior draws can then be obtained by sequentially sampling from: 1) $p(A, \Sigma_r \mid Y, B, \Sigma_c, \kappa, \omega)$; 2) $p(B, \Sigma_c \mid Y, A, \Sigma_r, \kappa, \omega)$; 3) $p(\kappa \mid Y, A, B, \Sigma_r, \Sigma_c, p_o)$; 4) $p(\omega \mid Y, A, B, \Sigma_r, \Sigma_c, \kappa, p_o)$; and 5) $p(p_o \mid Y, A, B, \Sigma_r, \Sigma_c, \kappa, \omega)$. Steps 1-3 remain the same as before. To implement Step 4, we discretize the distribution using a fine grid as proposed in Stock and Watson (2016). Consequently, each $o_t$ follows a discrete distribution that can be easily sampled from. In particular, we have

$$p(\omega \mid Y, A, B, \Sigma_r, \Sigma_c, \kappa, p_o) = \prod_{t=1}^{T} p(o_t \mid Y, A, B, \Sigma_r, \Sigma_c, \kappa, p_o) \propto \prod_{t=1}^{T} p(o_t \mid p_o) o_t^{-nk} e^{-\frac{s_t^2}{2o_t}},$$

where $p(o_t \mid p_o)$ is the prior for $o_t$ and $s_t^2 = \text{tr} \left( \Sigma_{\omega}^{-1} \Sigma_{\epsilon}^{-1} \Sigma_{\epsilon} \right)$. Hence, we can sample each $\omega_t$ from its discrete distribution. Lastly, Step 5 can be implemented easily as $p_o$ follows
the following beta distribution

\[ (p_o \mid Y, A, B, \Sigma_r, \Sigma_c, \kappa, \omega) \sim B(a_0 + n_o, b_0 + T - n_o), \]

where \( n_o = \sum_{t=1}^{T} 1(o_t > 1) \) is the number of outliers.

**Example 3 The common stochastic volatility**

Next, we incorporate the common stochastic volatility introduced in Carriero, Clark, and Marcellino (2016) to our matrix autoregression with \( \omega_t = e^{h_t} \), where \( h_t \) follows an AR(1) process:

\[ h_t = \phi h_{t-1} + \varepsilon_t^h, \]

where \( \varepsilon_t^h \sim \mathcal{N}(0, \sigma^2_h) \).

We assume independent truncated normal and inverse-gamma priors for \( \phi \) and \( \sigma^2_h \): \( \phi \sim \mathcal{N}(\phi_0, V_\phi)1(\phi < 1) \) and \( \sigma^2_h \sim \mathcal{IG}(\nu_h, S_h) \). Then, posterior draws can be obtained by sampling from:

1) \( p(A, \Sigma_r \mid Y, B, \Sigma_c, \kappa, \omega) \);
2) \( p(B, \Sigma_c \mid Y, A, \Sigma_r, \kappa, \omega) \);
3) \( p(\kappa \mid Y, A, B, \Sigma_r, \Sigma_c, \omega, \phi, \sigma^2_h) \);
4) \( p(\omega \mid Y, A, B, \Sigma_r, \Sigma_c, \kappa, \phi, \sigma^2_h) \);
5) \( p(\phi \mid Y, A, B, \Sigma_r, \Sigma_c, \kappa, \omega, \sigma^2_h) \); and
6) \( p(\sigma^2_h \mid Y, A, B, \Sigma_r, \Sigma_c, \kappa, \omega, \phi) \).

Steps 1-2 again can be implemented exactly as described in Section 3 of the main text and Step 3 as outlined in Section A2. For Step 4, note that

\[
p(\omega \mid Y, A, B, \Sigma_r, \Sigma_c, \kappa, \phi, \sigma^2_h) = p(h \mid Y, A, B, \Sigma_r, \Sigma_c, \kappa, \phi, \sigma^2_h) = \prod_{t=1}^{T} p(Y_t \mid A, B, \Sigma_r, \Sigma_c, h_t),
\]

where \( p(h \mid \phi, \sigma^2_h) \) is a Gaussian density implied by the state equation,

\[
\log p(Y_t \mid A, B, \Sigma_r, \Sigma_c, h_t) = c_t - \frac{nk}{2}h_t - \frac{1}{2}e^{-h_t} s_t^2,
\]

and \( c_t \) is a normalizing constant that does not dependent on \( h_t \) and \( s_t^2 = \text{tr} \left( \Sigma^{-1}_c \mathbf{E}_t' \Sigma^{-1}_r \mathbf{E}_t \right) \).

It is easy to check that

\[
\frac{\partial}{\partial h_t} \log p(Y_t \mid A, B, \Sigma_r, \Sigma_c, h_t) = -\frac{nk}{2} + \frac{1}{2}e^{-h_t} s_t^2,
\]

\[
\frac{\partial^2}{\partial h_t^2} \log p(Y_t \mid A, B, \Sigma_r, \Sigma_c, h_t) = -\frac{1}{2}e^{-h_t} s_t^2.
\]

Then, one can implement a Newton-Raphson algorithm to obtain the mode of the log
density $\log p(h \mid Y, A, B, \Sigma_r, \Sigma_c, \phi, \sigma^2_h)$ and the negative Hessian evaluated at the mode, which are denoted as $\hat{h}$ and $K_h$, respectively. Using $\mathcal{N}(\hat{h}, K_h^{-1})$ as a proposal distribution, one can sample $h$ directly using an acceptance-rejection Metropolis-Hastings step. We refer the readers to Chan (2017) for details. Finally, Steps 4 and 5 are standard and can be easily implemented (see., e.g., Chan and Hsiao, 2014).
Appendix B: Data

The US state-level data are sourced from the Federal Reserve Bank of St. Louis and the Bureau of Labor Statistics. For each of the 50 states, 6 quarterly time-series from 1991Q1 to 2023Q1 are obtained. Table 1 lists the 6 quarterly variables and describes how they are transformed. For example, $\Delta \log$ is used to denote the first difference in the logs, i.e.,

\[
\Delta \log y_t = \log y_t - \log y_{t-1}.
\]

<table>
<thead>
<tr>
<th>Variable</th>
<th>Source</th>
<th>Transformation</th>
</tr>
</thead>
<tbody>
<tr>
<td>Initial unemployment insurance claims</td>
<td>FRED</td>
<td>log</td>
</tr>
<tr>
<td>Continued unemployment insurance claims</td>
<td>FRED</td>
<td>log</td>
</tr>
<tr>
<td>Total nonfarm employment</td>
<td>BLS</td>
<td>400$\Delta \log$</td>
</tr>
<tr>
<td>Unemployment rate</td>
<td>FRED</td>
<td>no transformation</td>
</tr>
<tr>
<td>New housing permits</td>
<td>FRED</td>
<td>log</td>
</tr>
<tr>
<td>Real home price index</td>
<td>FRED</td>
<td>400$\Delta \log$</td>
</tr>
</tbody>
</table>

Table 1: Description of state-level variables in the empirical application.
References


