# Penalized estimation of sparse Markov regime-switching vector auto-regressive models

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#### Abstract

We consider sparse Markov regime-switching vector autoregressive (MSVAR) models in which the regimes are governed by a latent homogeneous Markov chain. In practice, even for moderate values of the number of Markovian regimes and data dimension, the associated MSVAR model has a large parameter dimension compared to a typical sample size. We provide a unified penalized conditional likelihood approach for estimating sparse MSVAR models. We show that our proposed estimators are consistent and recover the sparse structure of the model. We also show that, when the number of regimes is correctly or over-specified, our method provides consistent estimation of the predictive density. We develop an efficient implementation of the method based on a modified expectation-maximization (EM) algorithm. We discuss strategies for estimation of the number of regimes. We evaluate finite-sample performance of the method via simulations, and further demonstrate its utility by analyzing a real dataset. Supplementary materials for this paper are available online.

Keywords: EM algorithm, regularization methods, multivariate time series.

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### 1 Introduction

Markov regime-switching vector auto-regressive (MSVAR) models are ubiquitous in modelling heterogeneous and complex relationships between the variables of interest in multivariate time series analysis (Krolzig, 1997). The two stochastic components of an MSVAR are the observable time series vector  $\mathbf{Y}_t \in \mathbb{R}^d$ , and the latent Markov chain  $S_t \in \{1, \dots, M\}$ , for some finite  $M \in \mathbb{N}$ . The process  $S_t$  determines the VAR regime under which the conditional distribution of  $\mathbf{Y}_t | (S_t = s_t)$  evolves as a function of lagged  $\mathbf{Y}_{t-j}, j = 1, \dots, p \in \mathbb{N}$ . In this work, we focus on Gaussian MSVARs where the noise component in each VAR regime follows a zero-mean, d-dimensional Gaussian distribution. These models are applicable in a wide range of disciplines. For example, in macroeconomic time series such as manufacturing activities, consumer price indices, and housing and asset prices  $(\mathbf{Y}_t)$ , we typically see the effect of business cycles and the volatility clustering effect, which can be captured by  $S_t$  (Hamilton, 1988; Kim et al., 1998; Ang and Timmermann, 2012). In time series data related to natural phenomena such as wind power generation  $(\mathbf{Y}_t)$  across geographic regions, there are periods of high and low measurements that depend on wind speed  $(S_t)$  (Pinson and Madsen, 2012); similarly, there are periods of high and low temperature measurements that depend on annual meteorological cycles  $(S_t)$  corresponding to different seasons with warm and cold months (Monbet and Ailliot, 2017).

Maximum likelihood estimation (MLE) is the most common frequentist method of inference in MSVAR models. However, a limitation often encountered with MLE is the potentially large number of parameters to be estimated. In an MSVAR model with Mregimes and autoregressive order p, the total number of parameters is  $K = M(d + pd^2 + pd^2)$  d(d+1)/2 + M(M-1), which can be large even for moderate values of (d, p, M), compared to a typical sample size. For instance, in our case study each observation is 10-dimensional, and for an MSVAR model with AR-order p = 1 and the number of regimes M ranging from 1 to 5, there are  $165 \le K \le 845$  parameters to estimate based on a sample of size 481. Therefore, besides an obfuscated model interpretation, it can also be difficult to perform stable MLE in large-dimensional parameter spaces. It thus becomes essential to consider strategies that enable more stable and interpretable parameter estimation. With this motivation, we perform parameter estimation using regularization techniques that have been successful in both high-dimensional VAR and covariance estimation problems (Basu and Matteson, 2021; Lam, 2020). These techniques arise from the assumption that many of the model parameters are null. In the context of MSVAR models, we assume that both VAR coefficient matrices and the noise covariance–or precision–matrices are sparse, that is, many of their entries are zero. This also results in more meaningful model interpretations.

There has been a huge surge in research on estimation in sparse VAR models (Basu and Michailidis, 2015; Nicholson et al., 2020, and references therein), and sparse covariance and precision matrix estimation (Friedman et al., 2008; Shojaie and Michailidis, 2010; Bien and Tibshirani, 2011; Lam, 2020, and references therein). This has been led by the development of penalized methods such as the LASSO (Tibshirani, 1996), adaptive LASSO (Zou, 2006), SCAD (Fan and Li, 2001) and MCP (Zhang, 2010). However, to the best of our knowledge, there are only a few works on regularized estimation for MSVARs. In the context of hidden Markov models (HMMs)–which are MSVARs with AR-order p = 0–Städler and Mukherjee (2013) studied the graphical LASSO algorithm for estimation of sparse precision matrices in high-dimensions. Fiecas et al. (2017) proposed a shrinkage, rather than sparsity, approach to obtain stable covariance matrix estimates in high-dimensional HMMs. To simulate temperature data using a sparse MSVAR with a non-homogeneous Markov chain, Monbet and Ailliot (2017) performed simultaneous VAR coefficient and precision matrix estimation.

In this work, we focus on a general sparse estimation framework for MSVAR models, which is not available in the reviewed literature. We propose a penalized conditional likelihood approach that allows for sparse estimation of the regime-specific VAR coefficients and covariance/precision matrices. We condition on the initial state of the Markov chain  $S_t$ , which avoids the estimation of its initial distribution and thus simplifies the problem. We show that, irrespective of the initial state conditioning, our method consistently recovers the sparse MSVAR model. We also provide an estimator for the *h*-step-ahead predictive density, and show that as long as the true number of AR regimes is not under-specified, this estimator is consistent for the true predictive density. All of our theoretical results are provided in the Supplement, Section A2. We implement the proposed method by using a modified EM algorithm, combined with a generalized gradient descent method in the Mstep. We investigate strategies for estimation of the number of AR regimes. We study finitesample performance of the proposed methods via simulations, and further demonstrate their utility by analyzing a Canadian macroeconomic dataset.

Section 2 presents sparse MSVAR models. Section 3 describes the proposed estimation methods and their numerical implementation. We discuss the computation of predictive densities in Section 4. Section 5 presents a simulation study, and Section 6 contains a case study on a Canadian macroeconomic dataset to demonstrate the usage of our methodology.

### 2 Sparse Markov regime-switching vector auto-regressives

Let  $\{S_t \in \{1, \ldots, M\}, t = 1, 2, \ldots\}$  be a latent homogeneous Markov chain for some finite  $M \in \mathbb{N}$ . We denote its  $M \times M$  transition probability matrix by P with the (i, j)-th entry

$$\mathbb{P}(S_t = j | S_{t-1} = i, S_{t-2} = s_{t-2}, \dots, S_1 = s_1) = \mathbb{P}(S_t = j | S_{t-1} = i) = \alpha_{ij},$$

which is the probability of entering state j at time t from state i at time t - 1, and  $\sum_{j=1}^{M} \alpha_{ij} = 1, i = 1, \ldots, M$ . Further, let  $\{\mathbf{Y}_t \in \mathbb{R}^d, t = 1, 2, \ldots\}$  be an observable time series with observed values  $\{\mathbf{y}_t, t = 1, 2, \ldots\}$ . For any t > p, we assume that conditional on  $\mathbf{Y}_{1:t-1} = \mathbf{y}_{1:t-1}$  and  $S_{p:n} = s_{p:n}$ , the distribution of  $\mathbf{Y}_t$  only depends on the lagged  $\mathbf{Y}_{t-p:t-1} = \mathbf{y}_{t-p:t-1}$  and  $S_t = s_t$ . In a Gaussian MSVAR with M regimes and AR-order p, the conditional distribution of  $\mathbf{Y}_t$  given  $S_t = m$  and  $\mathbf{Y}_{t-j} = \mathbf{y}_{t-j}, j = 1, \ldots, p$ , is a state-dependent multivariate Gaussian with covariance matrix  $\mathbf{\Sigma}^{(m)} = (\sigma_{ij}^{(m)}) \in \mathbb{R}^{d \times d}$  and time-dependent mean vector

$$\boldsymbol{\mu}_{t}^{(m)} := \boldsymbol{\nu}^{(m)} + \boldsymbol{A}_{1}^{(m)} \mathbf{y}_{t-1} + \ldots + \boldsymbol{A}_{p}^{(m)} \mathbf{y}_{t-p}, \tag{1}$$

where  $\boldsymbol{\nu}^{(m)} = (\nu_1^{(m)}, \dots, \nu_d^{(m)})^{\mathsf{T}} \in \mathbb{R}^d$ , and  $\boldsymbol{A}_l^{(m)} = (a_{l,ij}^{(m)}) \in \mathbb{R}^{d \times d}, l = 1, \dots, p$ , are the AR coefficient matrices. In simple terms, the behaviour of  $\mathbf{Y}_t$  within each regime-determined by the latent process  $S_t$ -over time is modeled by a Gaussian VAR with AR-order p.

The parameters of interest are the transition probability matrix  $\boldsymbol{P}$ , the AR intercepts  $\boldsymbol{\nu}^{(1:M)} := \operatorname{vec}([\boldsymbol{\nu}^{(1)}, \dots, \boldsymbol{\nu}^{(M)}])$ , the AR coefficients  $\boldsymbol{A}^{(1:M)} := \operatorname{vec}([\boldsymbol{A}^{(1)}, \dots, \boldsymbol{A}^{(M)}])$ , with  $\boldsymbol{A}^{(m)} := \operatorname{vec}([\boldsymbol{A}^{(m)}_1, \dots, \boldsymbol{A}^{(m)}_p])$ ,  $m = 1, \dots, M$ , the covariance matrices  $\boldsymbol{\Sigma}^{(1:M)} := \operatorname{vec}([\boldsymbol{\Sigma}^{(1)}, \dots, \boldsymbol{\Sigma}^{(M)}])$ , and the precision matrices  $\boldsymbol{\Omega}^{(1:M)} := \operatorname{vec}([\boldsymbol{\Omega}^{(1)}, \dots, \boldsymbol{\Omega}^{(M)}])$ , where  $\boldsymbol{\Omega}^{(m)} = (\boldsymbol{\Sigma}^{(m)})^{-1} = (\omega_{ij}^{(m)})$ . Let  $\boldsymbol{\theta} \in \boldsymbol{\Theta} \subseteq \mathbb{R}^K$  be the vector of all parameters with

 $K = M(d + pd^2 + d(d + 1)/2) + M(M - 1)$ . In our estimation procedure, we directly estimate either  $\Sigma^{(m)}$  or  $\Omega^{(m)}$ , and the parameter  $\theta$  includes either  $\Sigma^{(1:M)}$  or  $\Omega^{(1:M)}$ .

**Conditional likelihood:** Let  $\mathbf{y}_{1:n} := {\mathbf{y}_1, \dots, \mathbf{y}_n}$  be observations from a Gaussian MSVAR model with M regimes and AR-order p. We consider the joint conditional density of  $\mathbf{Y}_{p+1:n}$  given  $(\mathbf{y}_{1:p}, S_p = s_p)$ . By the total probability rule, we have

$$f_{\boldsymbol{\theta}}(\mathbf{y}_{p+1:n}|\mathbf{y}_{1:p}, s_p) = \sum_{s_n=1}^M \dots \sum_{s_{p+1}=1}^M f(\mathbf{y}_{p+1:n}|\mathbf{y}_{1:p}, S_{p:n} = s_{p:n}) \mathbb{P}(S_{p+1:n} = s_{p+1:n}|\mathbf{y}_{1:p}, s_p).$$
(2)

By the homogeneity of  $S_t$ , conditional independence of  $\mathbf{Y}_t$ 's, and the Gaussianity, we obtain

$$\begin{split} \mathbb{P}(S_{p+1:n} = s_{p+1:n} | \mathbf{y}_{1:p}, S_p = s_p) &= \mathbb{P}(S_{p+1:n} = s_{p+1:n} | S_p = s_p) = \prod_{t=p+1}^n \alpha_{s_{t-1}, s_t}, \\ f(\mathbf{y}_{p+1:n} | \mathbf{y}_{1:p}, S_{p:n} = s_{p:n}) &= \prod_{t=p+1}^n f(\mathbf{y}_t | \mathbf{y}_{1:t-1}, S_{p:n} = s_{p:n}) = \prod_{t=p+1}^n f(\mathbf{y}_t | \mathbf{y}_{t-p:t-1}, S_t = s_t) \\ &= \prod_{t=p+1}^n \phi(\mathbf{y}_t; \boldsymbol{\mu}_t^{(s_t)}, \boldsymbol{\Sigma}^{(s_t)}), \end{split}$$

where  $\phi(\cdot; \boldsymbol{\mu}, \boldsymbol{\Sigma})$  is the *d*-dimensional Gaussian density. Therefore, the density (2) becomes

$$f_{\theta}(\mathbf{y}_{p+1:n}|\mathbf{y}_{1:p}, s_p) = \sum_{s_n=1}^{M} \dots \sum_{s_{p+1}=1}^{M} \left(\prod_{t=p+1}^{n} \alpha_{s_{t-1}, s_t}\right) \left(\prod_{t=p+1}^{n} \phi(\mathbf{y}_t; \boldsymbol{\mu}_t^{(s_t)}, \boldsymbol{\Sigma}^{(s_t)})\right),$$

where  $\boldsymbol{\mu}_{t}^{(s_{t})}$  is given in (1). The conditioning on  $\mathbf{y}_{1:p}$  is standard in time series when fitting AR-type models. We also condition on  $S_{p} = s_{p}$  to avoid specification of the initial distribution of  $S_{t}$  which is an intricate issue in itself. The effect of conditioning on  $S_{p}$  dissipates asymptotically (Douc et al., 2004). Thus, for any fixed  $s_{p}$ , the conditional log-likelihood is

$$l_n(\boldsymbol{\theta}; s_p) := \log \left( f_{\boldsymbol{\theta}} \big( \mathbf{y}_{p+1:n} | \mathbf{y}_{1:p}, s_p \big) \right).$$
(3)

One could obtain the (conditional) MLE of  $\boldsymbol{\theta}$  by maximizing (3) with respect to  $\boldsymbol{\theta}$ . However, as discussed in the introduction, even for moderate values of (d, p, M), the parameter  $\boldsymbol{\theta}$  has a large dimension which can result in an unstable estimation. Hence, to attain a more stable and meaningful estimation of  $\boldsymbol{\theta}$ , we assume that the data-generating model is sparse so that the number of parameters to estimate is much less than K. We develop an estimation framework based on penalization of the conditional likelihood (3) in Section 3. **Sparsity:** Different sparsity structures on the AR coefficients  $\{\boldsymbol{A}_{l}^{(m)}, m = 1, \ldots, M, l = 1, \ldots, p\}$  can be assumed for specific applications, for example, those in Nicholson et al. (2020). For generality, we do not assume any specific sparsity structure. Nonetheless, our method can be adapted to obtain structured sparse AR coefficients' estimates by an appropriate modification of the penalty function in Section 3.

Estimation of the covariance  $\Sigma^{(m)}$  or precision  $\Omega^{(m)}$  matrices when the dimension d is large is even more challenging, as their MLE estimates can be particularly unstable. To circumvent this, one could either assume a sparse structure on these matrices, or use a shrinkage-based approach to estimate them (Fiecas et al., 2017). We focus on the former approach, and assume that the aforementioned matrices have many off-diagonal zero entries. The sparsity assumption on covariance or precision matrices is application-dependent (Friedman et al., 2008; Bien and Tibshirani, 2011), and we study both scenarios separately. Figure 5 illustrates an example of a sparse 3-regime MSVAR model for macroeconomic variables and the estimated sparse MSVAR parameters and regimes (Section 6).

### 3 Maximum penalized likelihood estimation

We estimate  $\theta$  by penalizing the conditional log-likelihood in (3) using an appropriate penalty function. This results in more stable parameter estimates. Specifically, we obtain the maximum penalized conditional likelihood estimate (MPLE) of  $\boldsymbol{\theta}$  as

$$\widehat{\boldsymbol{\theta}}_{n} \in \arg \max \left\{ \mathcal{L}_{n}(\boldsymbol{\theta}; s_{p}) := \frac{1}{n-p} l_{n}(\boldsymbol{\theta}; s_{p}) - R(\boldsymbol{\theta}; \lambda_{1}, \lambda_{2}) \right\},$$
(4)  
with 
$$R(\boldsymbol{\theta}; \lambda_{1}, \lambda_{2}) := \sum_{m=1}^{M} \sum_{l=1}^{p} \sum_{i,j=1}^{d} r_{\lambda_{1}}(a_{l,ij}^{(m)}) + \sum_{m=1}^{M} \sum_{\substack{i,j=1\\j\neq i}}^{d} r_{\lambda_{2}}(\gamma_{ij}^{(m)}),$$

where  $a_{l,ij}^{(m)}$  are the AR coefficients, and  $\gamma_{ij}^{(m)}$  correspond to either  $\sigma_{ij}^{(m)}$  or  $\omega_{ij}^{(m)}$ . The  $r_{\lambda}$ is a penalty function with tuning parameter  $\lambda > 0$  that determines the severity of the penalty and consequently the level of sparsity of  $\hat{\theta}_n$ . In our numerical studies, we use the well-known penalties LASSO, adaptive LASSO (ADALASSO), MCP and SCAD, which are given in the Supplement, Section A1.1. General assumptions on the choice of  $r_{\lambda}$  are given in the Supplement, Section A2. By design, many entries of  $\hat{\theta}_n$  corresponding to  $\hat{A}^{(1:M)}$  and  $\hat{\Sigma}^{(1:M)}$ or  $\hat{\Omega}^{(1:M)}$  are zero, resulting in an estimated sparse MSVAR. Theorem 1 (Supplement, Section A2) states that under the correct specification of M and appropriate choices of  $(r_{\lambda}, \lambda)$ , there exists a local maximizer  $\hat{\theta}_n$  of the penalized conditional likelihood  $\mathcal{L}_n(\theta; s_p)$  that is a consistent and sparse estimator of the true parameter  $\theta^*$ . As there is no closed-form solution to the optimization in (4), we provide a numerical algorithm to approximate  $\hat{\theta}_n$ .

### 3.1 Modified EM algorithm

The EM algorithm is commonly used to perform MLE in a wide range of latent variable models, including MSVAR (Krolzig, 1997). The main advantage of this algorithm compared to a direct gradient ascent method for maximizing  $\mathcal{L}_n(\boldsymbol{\theta}; s_p)$  is its ease of implementation in the presence of the latent regime-governing  $S_t$ . Nevertheless, we use gradient ascent method in the maximization (M-) step of the EM algorithm for obtaining the estimates of AR coefficient and covariance/precision matrices. We adapt the EM algorithm to our penalization method by first introducing a so-called penalized complete-data likelihood.

For each  $S_t$ , we introduce a vector  $\boldsymbol{\xi}_t := (\xi_{t1}, \dots, \xi_{tM})^{\mathsf{T}}$ , where  $\xi_{ti} = \mathbb{1}_{\{S_t=i\}}$  represents the membership of  $\mathbf{y}_t$  to regime *i*, so that  $\sum_{i=1}^M \xi_{ti} = 1$ . We denote the complete data as  $\{(\mathbf{y}_t, \boldsymbol{\xi}_t), t = p + 1, \dots, n\}$ , and the penalized complete conditional log-likelihood is

$$\mathcal{L}_{n}^{c}(\boldsymbol{\theta};s_{p}) := \sum_{i,j=1}^{M} \sum_{t=p+1}^{n} \frac{\xi_{(t-1)i}\xi_{tj}\log\alpha_{ij}}{n-p} + \sum_{i=1}^{M} \sum_{t=p+1}^{n} \frac{\xi_{ti}\log\phi(\mathbf{y}_{t};\boldsymbol{\mu}_{t}^{(i)},\boldsymbol{\Sigma}^{(i)})}{n-p} - R(\boldsymbol{\theta};\lambda_{1},\lambda_{2}).$$

Clearly, the function  $\mathcal{L}_n^c$  cannot be used directly to estimate  $\boldsymbol{\theta}$  since  $\boldsymbol{\xi}_t$  are non-observable. Instead, at the (k + 1)-th iteration, the algorithm proceeds in two steps as follows. **F** step: It computes the conditional expectation of  $\mathcal{L}_n^c$  with respect to  $\boldsymbol{\xi}_n$  given  $\boldsymbol{x}_n$  and

**E-step**: It computes the conditional expectation of  $\mathcal{L}_n^c$  with respect to  $\boldsymbol{\xi}_t$  given  $\mathbf{y}_{1:n}$  and the current update  $\boldsymbol{\theta}^{(k)}$ . This results in

$$Q(\boldsymbol{\theta}; \boldsymbol{\theta}^{(k)}, s_p) := \mathbb{E}(\mathcal{L}_n^c | \mathbf{y}_{1:n}, \boldsymbol{\theta}^{(k)}, s_p) = \frac{1}{n-p} \sum_{i,j=1}^M \sum_{t=p+1}^n \zeta_{t,ij}^{(k)} \log \alpha_{ij} - \frac{1}{n-p} \sum_{i=1}^M \sum_{t=p+1}^n \zeta_{ti}^{(k)} \left( \log |\boldsymbol{\Sigma}^{(i)}| + \frac{1}{2} (\mathbf{y}_t - \boldsymbol{\mu}_t^{(i)})^\mathsf{T} \boldsymbol{\Omega}^{(i)} (\mathbf{y}_t - \boldsymbol{\mu}_t^{(i)}) \right) - R(\boldsymbol{\theta}; \lambda_1, \lambda_2),$$

where, for  $t = p + 1, \ldots, n$  and  $i, j = 1, \ldots, M$ , the weights

$$\zeta_{t,ij}^{(k)} = \mathbb{P}(\xi_{(t-1)i} = 1, \xi_{tj} = 1 | \mathbf{y}_{1:n}, \boldsymbol{\theta}^{(k)}, s_p), \quad \zeta_{ti}^{(k)} = \mathbb{P}(\xi_{ti} = 1 | \mathbf{y}_{1:n}, \boldsymbol{\theta}^{(k)}, s_p), \quad (5)$$

are computed using a recursive forward-backward procedure as follows.

We define the vectors  $\boldsymbol{\eta}_t := \left(\phi(\mathbf{y}_t; \boldsymbol{\mu}_t^{(1)}; \boldsymbol{\Sigma}^{(1)}), \dots, \phi(\mathbf{y}_t; \boldsymbol{\mu}_t^{(M)}; \boldsymbol{\Sigma}^{(M)})\right)^{\mathsf{T}}$  and  $\widehat{\boldsymbol{\xi}}_{t|\tau} := \mathbb{E}(\boldsymbol{\xi}_t | \mathbf{y}_{1:\tau}) = \left(\mathbb{P}(\xi_{t1} = 1 | \mathbf{y}_{1:\tau}), \dots, \mathbb{P}(\xi_{tM} = 1 | \mathbf{y}_{1:\tau})\right)^{\mathsf{T}}$ , for  $p \leq \tau \leq t$  and  $t \geq p + 1$ . We first compute  $\widehat{\boldsymbol{\xi}}_{t|t}$ (forward recursion) and use them to obtain  $\widehat{\boldsymbol{\xi}}_{t|n}$  (backward recursion). By Bayes' rule,

$$\mathbb{P}(\xi_{tm} = 1 | \mathbf{y}_{1:t}) = \mathbb{P}(\xi_{tm} = 1 | \mathbf{y}_t, \mathbf{y}_{1:t-1}) = \frac{\mathbb{P}(\mathbf{y}_t | \xi_{tm} = 1, \mathbf{y}_{1:t-1}) \mathbb{P}(\xi_{tm} = 1 | \mathbf{y}_{1:t-1})}{\mathbb{P}(\mathbf{y}_t | \mathbf{y}_{1:t-1})}.$$
 (6)

The first term in the numerator of (6) is the *m*-th entry of  $\eta_t$ , m = 1, ..., M. For the second term in the numerator, using the total probability rule and the homogeneity assumption,

$$\mathbb{P}(\xi_{tm} = 1 | \mathbf{y}_{1:t-1}) = \sum_{i=1}^{M} \mathbb{P}(\xi_{tm} = 1 | \xi_{(t-1)i} = 1, \mathbf{y}_{1:t-1}) \mathbb{P}(\xi_{(t-1)i} = 1 | \mathbf{y}_{1:t-1})$$
$$= \sum_{i=1}^{M} \alpha_{im} \mathbb{P}(\xi_{(t-1)i} = 1 | \mathbf{y}_{1:t-1}), \ m = 1, \dots, M$$
(7)

or, in its vector form,  $\widehat{\boldsymbol{\xi}}_{t|t-1} = \boldsymbol{P}^{\mathsf{T}} \widehat{\boldsymbol{\xi}}_{t-1|t-1}$ . For the denominator in (6), we thus have

$$\mathbb{P}(\mathbf{y}_t | \mathbf{y}_{1:t-1}) = \sum_{m=1}^{M} \mathbb{P}(\mathbf{y}_t | \xi_{(t-1)m} = 1, \mathbf{y}_{1:t-1}) \mathbb{P}(\xi_{(t-1)m} = 1 | \mathbf{y}_{1:t-1})$$
$$= \boldsymbol{\eta}_t^{\mathsf{T}} \boldsymbol{P}^{\mathsf{T}} \widehat{\boldsymbol{\xi}}_{t-1|t-1} = \boldsymbol{\eta}_t^{\mathsf{T}} \widehat{\boldsymbol{\xi}}_{t|t-1}.$$
(8)

Using the definition of  $\eta_t$ , (7) and (8), we then compute (6). Next, we compute the elements of  $\hat{\boldsymbol{\xi}}_{t|n}$  as follows. By Bayes' rule and the model assumptions, for  $m = 1, \ldots, M$ ,

$$\mathbb{P}(\xi_{tm} = 1 | \mathbf{y}_{1:n}) = \sum_{i=1}^{M} \mathbb{P}(\xi_{tm} = 1, \xi_{(t+1)i} = 1 | \mathbf{y}_{1:n}) \\
= \sum_{i=1}^{M} \frac{\alpha_{mi} \ \mathbb{P}(\xi_{tm} = 1 | \mathbf{y}_{1:t}) \mathbb{P}(\xi_{(t+1)i} = 1 | \mathbf{y}_{1:n})}{\mathbb{P}(\xi_{(t+1)i} = 1 | \mathbf{y}_{1:t})},$$
(9)

which, as shown in Section A1.2 of the Supplement, uses that

$$\mathbb{P}(\xi_{tm} = 1, \xi_{(t+1)i} = 1 | \mathbf{y}_{1:n}) = \frac{\mathbb{P}(\xi_{(t+1)i} = 1 | \xi_{tm} = 1) \mathbb{P}(\xi_{tm} = 1 | \mathbf{y}_{1:t}) \mathbb{P}(\xi_{(t+1)i} = 1 | \mathbf{y}_{1:n})}{\mathbb{P}(\xi_{(t+1)i} = 1 | \mathbf{y}_{1:t})}.$$
(10)

Letting  $\odot$  and  $\oslash$  be the element-wise vector product and division, respectively, we write (6) and (9) in their vector forms,

• Forward recursion (filtering): for t = p + 1, ..., n,

$$\widehat{\boldsymbol{\xi}}_{t|t} = \frac{\boldsymbol{\eta}_t \odot \widehat{\boldsymbol{\xi}}_{t|t-1}}{\mathbf{1}_M^{\mathsf{T}} \left( \boldsymbol{\eta}_t \odot \widehat{\boldsymbol{\xi}}_{t|t-1} \right)} = \frac{\boldsymbol{\eta}_t \odot \boldsymbol{P}^{(k)} \widehat{\boldsymbol{\xi}}_{t-1|t-1}}{\mathbf{1}_M^{\mathsf{T}} \left( \boldsymbol{\eta}_t \odot \boldsymbol{P}^{(k)} \widehat{\boldsymbol{\xi}}_{t-1|t-1} \right)}.$$
(11)

• Backward recursion (smoothing): for  $t = n - 1, \dots, p + 1$ ,

$$\widehat{\boldsymbol{\xi}}_{t|n} = \left[ \boldsymbol{P}^{(k)^{\mathsf{T}}} \left( \widehat{\boldsymbol{\xi}}_{t+1|n} \oslash \widehat{\boldsymbol{\xi}}_{t+1|t} \right) \right] \odot \widehat{\boldsymbol{\xi}}_{t|t}.$$
(12)

Finally, we set  $\zeta_{ti}^{(k)} = \mathbb{P}(\xi_{ti} = 1 | \mathbf{y}_{1:n}, \boldsymbol{\theta}^{(k)}, s_p)$  in (5) as the *i*-th entry of the vector  $\widehat{\boldsymbol{\xi}}_{t|n}$ , for  $i = 1, \ldots, M$ . The joint probabilities  $\zeta_{t,ij}^{(k)} = \mathbb{P}(\xi_{(t-1)i} = 1, \xi_{tj} = 1 | \mathbf{y}_{1:n}, \boldsymbol{\theta}^{(k)}, s_p)$  in (5), for  $i, j = 1, \ldots, M$ , are the ((j-1)M + i)-th entries of the vector

$$\widehat{\boldsymbol{\xi}}_{t|n}^{\text{joint}} := (\zeta_{t,11}^{(k)}, \dots, \zeta_{t,MM}^{(k)})^{\mathsf{T}} = \operatorname{vec}(\boldsymbol{P}^{(k)}) \odot \left[ \left( \widehat{\boldsymbol{\xi}}_{t|n} \oslash \widehat{\boldsymbol{\xi}}_{t|t-1} \right) \otimes \widehat{\boldsymbol{\xi}}_{t-1|t-1} \right],$$
(13)

which are computed using (10), for t = p+1, ..., n. Here,  $\otimes$  denotes the Kronecker product. The details of all the above derivations are given in Section A1.2 of the Supplement. **M-step**: It maximizes Q with respect to  $\boldsymbol{\theta}$ , resulting in the updates  $\boldsymbol{\theta}^{(k+1)}$ . Due to the structure of Q, we separately maximize it with respect to  $\boldsymbol{P}$ ,  $\{\boldsymbol{\nu}^{(m)}\}_{m=1}^{M}$ , and  $\{\boldsymbol{A}^{(m)}, \boldsymbol{\Sigma}^{(m)}\}_{m=1}^{M}$ .

The updates for the transition probabilities  $\alpha_{ij}$  and the intercepts  $\boldsymbol{\nu}^{(m)}$  are given by

$$\alpha_{s_{p},j}^{(k+1)} = \frac{\sum_{t=p+1}^{n} \zeta_{t,s_{p},j}^{(k)}}{\sum_{l=1}^{M} \sum_{t=p+1}^{n} \zeta_{t,s_{p},l}^{(k)}}, \quad \alpha_{ij}^{(k+1)} = \frac{\sum_{t=p+2}^{n} \zeta_{t,ij}^{(k)}}{\sum_{l=1}^{M} \sum_{t=p+2}^{n} \zeta_{t,il}^{(k)}}, \quad i, j = 1..., M, \ i \neq s_{p}; 14)$$
$$\boldsymbol{\nu}^{(m,k+1)} = \frac{\sum_{t=p+1}^{n} \zeta_{tm}^{(k)} (\mathbf{y}_{t} - \sum_{i=1}^{p} \boldsymbol{A}_{i}^{(m,k)} \mathbf{y}_{t-i})}{\sum_{t=p+1}^{n} \zeta_{tm}^{(k)}}, \quad m = 1, ..., M.$$
(15)

Regarding the AR coefficients, we first update  $A^{(m)}$ , and then  $\Sigma^{(m)}$  or  $\Omega^{(m)}$ , for each regime  $m = 1, \ldots, M$ . The optimization problems for these updates are, respectively,

$$\min_{\{\boldsymbol{A}_{l}^{(m)}\}_{l=1}^{p}} \frac{1}{2(n-p)} \sum_{t=p+1}^{n} \zeta_{tm}^{(k)} \big( \mathbf{y}_{t} - \bar{\boldsymbol{\mu}}_{t}^{(m)} \big)^{\mathsf{T}} \widehat{\boldsymbol{\Omega}}^{(m,k)} \big( \mathbf{y}_{t} - \bar{\boldsymbol{\mu}}_{t}^{(m)} \big) + \sum_{l=1}^{p} \sum_{i,j=1}^{d} r_{\lambda_{1}} \big( a_{l,ij}^{(m)} \big), \quad (16)$$

$$\min_{\boldsymbol{\Sigma}^{(m)} \succ 0} \frac{1}{2(n-p)} \Big( \widehat{n}_m \log |\boldsymbol{\Sigma}^{(m)}| + \operatorname{tr} \big( \boldsymbol{\Omega}^{(m)} \boldsymbol{S}^{(m)} \big) \Big) + \sum_{i \neq j=1}^d r_{\lambda_2} \big( \gamma_{ij}^{(m)} \big), \tag{17}$$

where  $\bar{\boldsymbol{\mu}}_{t}^{(m)}$  is (1) with  $\boldsymbol{\nu}^{(m)}$  replaced by  $\boldsymbol{\nu}^{(m,k+1)}$ . Further, we let  $\hat{n}_{m} := \sum_{t=p+1}^{n} \zeta_{tm}^{(k)}$ ,  $\boldsymbol{S}^{(m)} := \boldsymbol{U}^{\mathsf{T}} \boldsymbol{\Xi}^{(m)} \boldsymbol{U}$ , with  $\boldsymbol{\Xi}^{(m)} = \operatorname{diag} \{ \zeta_{p+1,m}^{(k)}, \dots, \zeta_{n,m}^{(k)} \}$ , and  $\boldsymbol{U} = (\mathbf{y}_{p+1} - \boldsymbol{\mu}_{p+1}^{(m,k+1)}, \dots, \mathbf{y}_{n} - \boldsymbol{\mu}_{n}^{(m,k+1)})^{\mathsf{T}}$ . The term  $\boldsymbol{\mu}_{t}^{(m,k+1)}$  is (1) with  $(\boldsymbol{\nu}^{(m)}, \boldsymbol{A}^{(m)})$  replaced by  $(\boldsymbol{\nu}^{(m,k+1)}, \boldsymbol{A}^{(m,k+1)})$ . We decouple the problem (16) coordinate-wise to alleviate the computation. Then both optimization problems in (16) and (17) can be written in a general form as  $\arg \min_{\mathbf{x}} \{f(\mathbf{x}) + R(\mathbf{x})\}$ , where f is differentiable and R is a penalty function. Using a generalized gradient descent, the solution is obtained via the updates

$$\mathbf{x}^{(j+1)} \in \arg\min_{\mathbf{x}} \frac{1}{2} \|\mathbf{x} - (\mathbf{z}^{(j)} - \frac{1}{u_{j+1}} \nabla f(\mathbf{z}^{(j)}))\|^2 + \frac{1}{u_{j+1}} R(\mathbf{x}),$$

where  $\mathbf{z}^{(j)}$  interpolates between  $\mathbf{x}^{(j)}$  and  $\mathbf{x}^{(j-1)}$ , and  $1/u_{j+1}$  is the descent step size. Since the penalty function R in (4) is decomposable, the updates have an analytical form for all the penalty functions we consider. Full details are given in the Supplement, Sections A1.3-A1.5, including an initialization of the EM algorithm and a data-dependent selection of tuning parameters  $(\lambda_1, \lambda_2)$ . Algorithm 1 summarizes the estimation steps.

Algorithm 1 Modified EM algorithm for maximum penalized likelihood estimation1: Initialize: k = 0,  $\alpha_{ij}^{(k)}$ ,  $\boldsymbol{\nu}^{(m,k)}$ ,  $\{\boldsymbol{A}_l^{(m,k)}\}_{l=1}^p$ ,  $\boldsymbol{\Sigma}^{(m,k)}$ , for  $i, j, m = 1, \dots, M$ ,  $\lambda_1, \lambda_2 \ge 0, \varepsilon > 0$ ;2: E-step: Compute  $\zeta_{ti}^{(k)}$ ,  $\zeta_{t,ij}^{(k)}$  in (5), for  $i, j = 1, \dots, M$ , using (11), (12), and (13);M-step: For  $i, j = 1, \dots, M$ , update  $\alpha_{ij}^{(k+1)}$  using (14);For  $m = 1, \dots, M$ , update  $\boldsymbol{\nu}^{(m,k+1)}$  using (15);For  $m = 1, \dots, M$ , update  $\{\boldsymbol{A}_l^{(m,k+1)}\}_{l=1}^p$  and  $\boldsymbol{\Sigma}^{(m,k+1)}$  or  $\boldsymbol{\Omega}^{(m,k+1)}$  solving (16)-(17);3: If  $\|\boldsymbol{\theta}^{(k)} - \boldsymbol{\theta}^{(k+1)}\| / \|\boldsymbol{\theta}^{(k)}\| < \varepsilon$ , return  $\boldsymbol{\theta}^{(k+1)}$ , else k = k + 1 and go to 2;

### **3.2** Selection of the number of regimes M

We use information-based criteria to estimate M when it is unknown. For a fitted MSVAR with M regimes and MPLE  $\hat{\theta}_n$ , we compute the regime-specific degrees of freedom as

$$D_m(\widehat{\theta}_n) = \sum_{l=1}^p \sum_{i,j=1}^d \mathbb{1}_{\{(\widehat{a}_{l,ij}^{(m)})\neq 0\}} + \sum_{i,j=1}^d \mathbb{1}_{\{\widehat{\gamma}_{ij}^{(m)}\neq 0\}}, \quad m = 1, \dots, M,$$

and let  $E_m(M) = M - 1 + D_m(\widehat{\theta}_n)$ . We denote the information criterion as

$$\mathcal{C}(M) = -2l_n(\widehat{\boldsymbol{\theta}}_n; s_p) + \sum_{m=1}^M c_m E_m(M), \qquad (18)$$

for some  $c_m > 0$ . The common choices for  $c_m$  are 2 (AIC) and  $\log(n-p)$  (BIC). Städler and Mukherjee (2013) introduced the mixture minimum description length (MDL) for HMMs by setting  $c_m = \log(\hat{n}_m)$ , where  $\hat{n}_m = \sum_{i=p+1}^n \hat{\zeta}_{im}$  is the estimated sample size for regime m and  $\hat{\zeta}_{im}$  are given in (5). We select the number of regimes over a set of candidate values  $\mathcal{M} = \{1, 2, \dots, M_{\max}\}$  as  $\widehat{M} \in \underset{M \in \mathcal{M}}{\operatorname{arg\,min}} \mathcal{C}(M)$ , for some pre-specified upper bound  $M_{\max} \in \mathbb{N}$ . Selection of the AR-order p is discussed in the Supplement, Section A1.6.

# 4 Prediction

Prediction is a highly relevant task in time series analysis. In stationary VAR models, an optimal prediction at time n + 1 given the observations  $\mathbf{y}_{1:n}$ , in terms of the minimum mean squared prediction error, is equal to the conditional expectation (Lütkepohl, 2007); h-step-ahead prediction is performed similarly. In this section, we compute the predictive density in MSVARs that can also be used to obtain the conditional expectations. For a fixed  $h \in \mathbb{N}$ , we compute the h-step-ahead predictive density  $f_M(\mathbf{y}_{n+1:n+h}|\mathbf{y}_{1:n})$  as follows. The one-step-ahead predictive density (h = 1) is

$$f_M(\mathbf{y}_{n+1}|\mathbf{y}_{1:n}) = \sum_{m=1}^M \mathbb{P}(S_{n+1} = m|\mathbf{y}_{1:n}) \phi(\mathbf{y}_{n+1}; \boldsymbol{\mu}_{n+1}^{(m)}, \boldsymbol{\Sigma}^{(m)}),$$
(19)

where the mean  $\boldsymbol{\mu}_{n+1}^{(m)}$  is given in (1), and the conditional probabilities  $\mathbb{P}(S_{n+1} = m | \mathbf{y}_{1:n}) = \sum_{j=1}^{M} \alpha_{jm} \mathbb{P}(S_n = j | \mathbf{y}_{1:n})$  can be readily computed using the forward-backward procedure (Supplement, Section A1.2). An initial conditional distribution  $\mathbb{P}(S_p = j | \mathbf{y}_{1:p})$  is also re-

quired to complete the computation. Since its effect on predictive density dissipates geometrically as n grows (Douc et al., 2004), and given the homogeneity of  $S_t$ , any non-informative distribution such as the discrete uniform  $\mathbb{P}(S_p = j | \mathbf{y}_{1:p}) = 1/M, j = 1, \dots, M$ , suffices.

Similarly, for  $h \ge 2$ , by the model assumptions, the predictive density is

$$f_{M}(\mathbf{y}_{n+1:n+h}|\mathbf{y}_{1:n}) = \sum_{m_{1:h}=1}^{M} \left\{ \mathbb{P}(S_{n+1}=m_{1},\ldots,S_{n+h}=m_{h}|\mathbf{y}_{1:n})f(\mathbf{y}_{n+1:n+h}|\mathbf{y}_{1:n},S_{n+1:n+h}=m_{1:h}) \right\}$$
$$= \sum_{m_{1:h}=1}^{M} \mathbb{P}(S_{n+1}=m_{1}|\mathbf{y}_{1:n}) \left(\prod_{t=2}^{h} \alpha_{m_{t-1},m_{t}}\right) \left(\prod_{t=1}^{h} \phi(\mathbf{y}_{n+t};\boldsymbol{\mu}_{n+t}^{(m_{t})},\boldsymbol{\Sigma}^{(m_{t})})\right) (20)$$

Given the MPLE  $\hat{\boldsymbol{\theta}}_n$  for a model with M regimes, the estimated predictive density  $\hat{f}_M(\mathbf{y}_{n+1:n+h}|\mathbf{y}_{1:n})$  is computed for  $h \ge 1$ , using either (19) or (20). The optimal h-step ahead predictor  $\hat{\mathbf{y}}_{n+h}$  in terms of minimum mean squared error is then the conditional expected value of  $\mathbf{Y}_{n+h}$ , given  $\mathbf{y}_{1:n}$ . For h = 1, the predicted value is

$$\widehat{\mathbf{y}}_{n+1} = \sum_{m=1}^{M} \widehat{\mathbb{P}}(S_{n+1} = m | \mathbf{y}_{1:n}) \left( \widehat{\boldsymbol{\nu}}^{(m)} + \sum_{l=1}^{p} \widehat{\boldsymbol{A}}_{l}^{(m)} \mathbf{y}_{n+1-l} \right)$$

For  $h \ge 2$ , predictions are obtained recursively by computing the predictions for  $\mathbf{y}_{n+j}$ ,  $j = 1, \ldots, h - 1$  (Lütkepohl, 2007; Krolzig, 1997). More specifically,

$$\widehat{\mathbf{y}}_{n+h} = \sum_{m=1}^{M} \widehat{\mathbb{P}}(S_{n+h} = m | \mathbf{y}_{1:n}) \left( \widehat{\boldsymbol{\nu}}^{(m)} + \sum_{l=1}^{p} \widehat{\boldsymbol{A}}_{l}^{(m)} \widetilde{\mathbf{y}}_{n+h-l} \right),$$

with  $\widetilde{\mathbf{y}}_{n+h-l} = \widehat{\mathbf{y}}_{n+h-l}$ , if (h-l) > 0 or  $\mathbf{y}_{n+h-l}$ , if  $(h-l) \le 0$ .

Theorem 2 (Supplement, Section A2) states that, if M is not under-specified, the estimated predictive density of the over-fitted model based on MPLE consistently estimates the *h*-step ahead predictive density (20) of the true MSVAR model. As a consequence, in practice when the true number of regimes is unknown, a rather conservative choice of Mconsidering the sample size n can guarantee a reliable estimate of (20), and of the optimal predictor in the sense of minimum mean squared prediction error.

### 5 Simulation study

Simulation design. We consider MSVAR models with p = 1, M = 2, 3, dimensions d = 20, 40, 100, and sample sizes n between 200 and 800. For these values of d, the parameter vector  $\theta^*$  of the data-generating MSVAR models has 104, 202 and 504 nonzero entries, respectively, and is designed as follows. We set the number of nonzero entries per matrix to be  $\lceil d/4 \rceil$ , randomly scattered on each matrix, and thus not assuming a specific sparsity structure in the AR and covariance or precision matrices. We set the true nonzero AR coefficients for each regime  $m = 1, \ldots, M$  as:  $\nu_i^{(m)*} \sim \mathcal{N}(m-2, 0.01), (a_l^{(m)*})_{ij} \sim \mathcal{N}((-1)^m, 0.01)$ . We ensure that the resulting MSVAR is stationary. Note that a VAR process with AR coefficients  $A_1, \ldots, A_p$  is stationary if the matrix

$$\begin{bmatrix} \boldsymbol{A}_1 \cdots \boldsymbol{A}_{p-1} & \boldsymbol{A}_p \\ \boldsymbol{I}_{d(p-1) \times d(p-1)} & \boldsymbol{0}_{d(p-1) \times d} \end{bmatrix}$$
(21)

has all its eigenvalues inside the unit circle (Lütkepohl, 2007, Section 9.1). In all of our simulations, to ensure stationarity of the VAR process for each Markov regime, we scale the AR matrices with entries  $(a_l^{(m)^*})_{ij}$  (generated as described above) to ensure that the modulus of each eigenvalue of the matrix in (21) is at most 0.75.

For the covariance matrices, we first sample  $\gamma_{ij}^{(m)} \sim \mathcal{N}(\frac{(-1)^m}{2}, 0.01), i \neq j, \gamma_{ii}^{(m)} \sim \mathcal{N}(0, 0.01)$ , and subsequently use one of the following formulas to obtain the nonzero elements, depending on the scenario, either covariance or precision matrices, under study:  $\sigma_{ij}^{(m)^*} = \frac{1}{2}(\gamma_{ij}^{(m)} + \gamma_{ji}^{(m)}), \ \omega_{ij}^{(m)^*} = \frac{1}{2}(\gamma_{ij}^{(m)} + \gamma_{ji}^{(m)})$ . To ensure positive-definiteness and numerical stability, if necessary, we shift the eigenvalues of matrices by adding a multiple of the identity matrix. This also aids in controlling the condition number (Rothman et al., 2008). We denote scenarios **S1** and **S2** respectively corresponding to sparse covariance and precision matrices, along with sparse AR coefficients.

The transition probability matrix  $\mathbf{P}^{\star}$  for M = 2 is set to:  $\alpha_{11}^{\star} = 0.75, \alpha_{22}^{\star} = 0.60$ ; and for M = 3 to:  $\alpha_{11}^{\star} = 0.5, \alpha_{12}^{\star} = 0.25, \alpha_{21}^{\star} = 0.2, \alpha_{22}^{\star} = 0.6, \alpha_{32}^{\star} = 0.25, \alpha_{33}^{\star} = 0.5$ . These choices test the estimation performance for an unequal number of data points in each regime.

**Performance measures.** Recall the MPLE  $\hat{\theta}_n \equiv \hat{\theta}$  of  $\theta^*$ , and let  $\tilde{\theta}_n \equiv \tilde{\theta}$  be the maximum likelihood estimate of  $\theta^*$  knowing its zero entries a priori. For a subvector  $\mathbf{v}$  of  $\theta^*$ , let  $\hat{\mathbf{v}}$  and  $\tilde{\mathbf{v}}$  be the corresponding subvectors of  $\hat{\theta}$  and  $\tilde{\theta}$ , respectively. Furthermore, let  $\widehat{M}$  be the estimate of the true number of regimes M using either BIC or MDL introduced in Section 3.2. We consider the following performance measures:

- relative estimation error:  $\operatorname{REE}(\widehat{\mathbf{v}}, \widetilde{\mathbf{v}}) = \operatorname{EE}(\widehat{\mathbf{v}}) / \operatorname{EE}(\widetilde{\mathbf{v}}) = \frac{||\widehat{\mathbf{v}} \mathbf{v}||}{||\mathbf{v}||} / \frac{||\widetilde{\mathbf{v}} \mathbf{v}||}{||\mathbf{v}||} \in [0, \infty);$
- true positive rate:  $\operatorname{TPR}(\widehat{\mathbf{v}}) = \frac{\sum_{i} \mathbbm{1}_{\{|\widehat{v}_i| > 0\}} \mathbbm{1}_{\{|v_i| > 0\}}}{\sum_{i} \mathbbm{1}_{\{|v_i| > 0\}}} \in [0, 1];$
- true negative rate:  $\text{TNR}(\widehat{\mathbf{v}}) = \frac{\sum_{i} \mathbbm{1}_{\{|\widehat{v}_i|=0\}} \mathbbm{1}_{\{|v_i|=0\}}}{\sum_{i} \mathbbm{1}_{\{|v_i|=0\}}} \in [0, 1];$
- correctly selected number of regimes:  $\mathbb{1}_{\{\widehat{M}=M\}} \in \{0,1\},\$

where  $|| \cdot ||$  represents the Euclidean norm and  $\mathbb{1}_{\{\cdot\}}$  is the indicator function. We compute EE, REE, TPR and TNR for different subvectors **v** corresponding to the AR coefficients and the covariance or precision matrices. We exclude the AR intercepts and the diagonals of the covariance or precision matrices when computing TPR and TNR.

Since in the computation of  $\tilde{\theta}$  we use the knowledge of the true zero parameters, in general, we expect  $\tilde{\theta}$  to outperform  $\hat{\theta}$  which estimates the zero parameters through penalization. Thus, we focus on the REE to quantify the performance of our estimation methodology relative to  $\tilde{\theta}$ , and seek to obtain the empirical mean or median of all the performance measures, except EE, close to 1. However, in the case of estimation of precision matrices, we observe REE < 1 which occurs due to the way we compute  $\tilde{\theta}$  without directly using the knowledge of the true zero parameters in the precision matrices. More details are provided in the Supplement, Section A3.

Our results are based on 50 simulated samples from each MSVAR model with a fixed parameter setting as described in the **Simulation design**. We start recording the generated observations at t = 401, to allow for a simulation burn-in. The first state is always set as  $s_{-399} = 1$ . For the estimation stage, we condition on  $s_p = 1$ . We analyze the results for scenario **S1** (sparse covariance matrix) here; those for scenario **S2** (sparse precision matrix) are given in the Supplement, Section A3, along with additional simulation designs on sparsity structure. We use BIC for selection of the tuning parameters ( $\lambda_1, \lambda_2$ ) using the method described in the Supplement, Section A1.5, as we find that AIC chooses over-parameterized models and therefore has a decreased performance.

The algorithms are implemented in C using the GNU Scientific Library (GSL) (Galassi et al., 2009) v2.7. The optimization procedure is implemented in parallel so that the estimation is done for all the regimes in an MSVAR model simultaneously.

### 5.1 Parameter estimation and sparsity

We fix M = 2, p = 1, and consider dimensions d = 20 with sample sizes  $n \in \{200, 300, 400\}$ , d = 40 with  $n \in \{300, 400, 500\}$ , and d = 100 with  $n \in \{600, 700, 800\}$ . For these values of d, the parameter vector  $\boldsymbol{\theta}$  to be estimated by our method has dimensions  $K = M(d + pd^2 + d(d+1)/2) + M(M-1) = 1262$ , 4922 and 30302, respectively. Thus, even though the change from d = 20 to 100 may appear moderate, the corresponding number of parameters to be estimated is very large. The results for d = 40, 100 are discussed below, and those for d = 20 are given in Section A3 of the Supplement; see also the end of this section.

Figure 1 shows the REE boxplots. We observe that for the AR coefficients, the ADALASSO and SCAD attain a performance close to  $\tilde{\theta}$ , followed by MCP and the LASSO. For covariance matrices, SCAD and MCP outperform the other penalties, and the medians reveal that they are comparable to  $\tilde{\theta}$ . Overall, SCAD and MCP attain a performance close to  $\tilde{\theta}$ , followed by the LASSO and ADALASSO. Estimation of covariance matrices is more difficult, as the medians depart from 1.0 more noticeably. In computing the overall REE, the estimation errors of the transition probabilities are included.

Figure 2 shows the TPR boxplots. We observe that the MPLE performs well under this measure. At least one penalty function attains an overall median TPR above 0.90 for the dimensions and sample sizes considered. The LASSO, SCAD and MCP perform similar to each other, while ADALASSO is less accurate. In terms of TNR (omitted), we find that for any setting, the median is at least 0.95 for both the AR and covariance matrices.

In summary, even though our theoretical results are proved in the setting of fixed dimension and large sample size, we conclude from the above experiments that our methodology has a good performance in finite-sample and moderate- to large-dimensions.

In Section A3 of the Supplement, we also provide the simulation results in terms of the EE and TPR for d = 20 and a wider range of sample sizes n to demonstrate the consistency and sparsity recovery property (Theorem 1) of the MPLE as n grows.

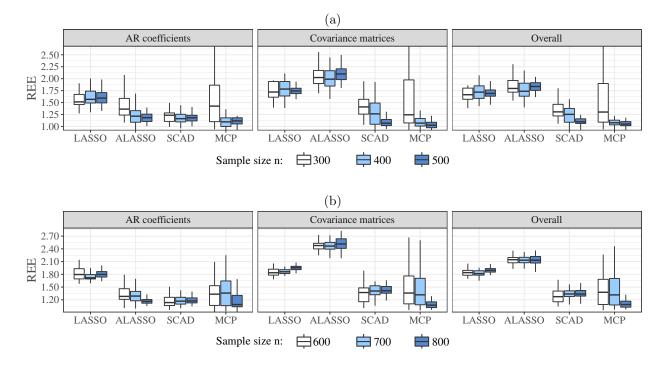


Figure 1: Relative estimation error (REE) based on 50 samples: (a) d = 40, K = 4922 (b) d = 100, K = 400, K

30302, where d and K are resp. the data and parameter dimensions.

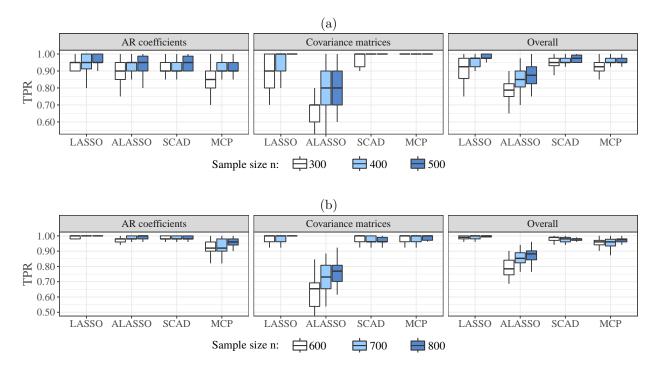


Figure 2: True positive rate (TPR, nonzero parameter detection) based on 50 random samples: (a) d = 40, K = 4922 (b) d = 100, K = 30302, where d and K are resp. the data and parameter dimensions.

### 5.2 Predictive density estimation and number of regime selection

We use models with  $M \in \{2, 3\}$ , p = 1, and generate samples of size n+h. We take d = 20, n = 800, and h = 8. We estimate the model parameters using n observations, setting the number of regimes M as either correctly or under/over-specified, ranging from 1 to 5. For the remaining h observations, we compute the estimated predictive densities in (20).

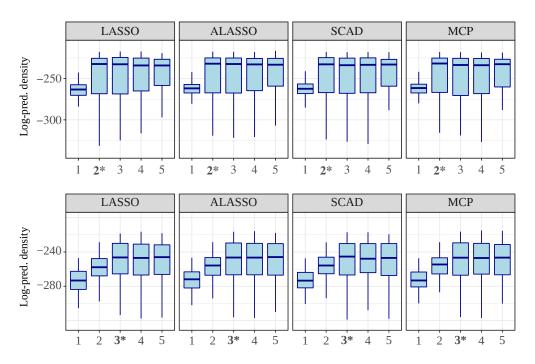


Figure 3: h-step-ahead log-predictive density (20) based on 50 random samples, for the true number of regimes  $M = 2^*$  (top),  $3^*$  (bottom), d = 20, n = 800, h = 8. The x-axis shows M values used in estimation.

Figure 3 shows boxplots of the estimated *h*-step ahead log-predictive density of  $\mathbf{y}_{n+1:n+h}$ , for different values of M. The results are based on 50 generated samples. We observe that the median values for the models with over-specified M fall within the inter-quantile range of the values for the model with correct M, which is expected as per the result of our Theorem 2 on the predictive density. The median for the under-specified M clearly deviates from the other medians, attaining a lower value. This is true for all the penalties. Next, we examine the performance of the criterion (18) for estimating M when the ARorder p is fixed. The results for estimating p are given in the Supplement, Section A1.6. We consider d = 20 and  $n \in \{400, 600, 800\}$ . We generate the data using models with true number of regimes M = 1, 2, 3, and estimate M using the BIC over the candidate values ranging from 1 to 5 for all the penalties. The mean of the performance measure  $1_{\{\widehat{M}=M\}}$ , with  $\widehat{M}$  being the estimated number of regimes, is reported in Figure 4. The ADALASSO outperforms other penalties and estimates the true M with a mean rate of at least 80% for any sample size. The performance for n = 600, 800 is very good with rates above 95%. The LASSO also appears more stable than SCAD and MCP. The performance of the MDL is not as accurate and omitted, which can be attributed to the small sample size per regime.

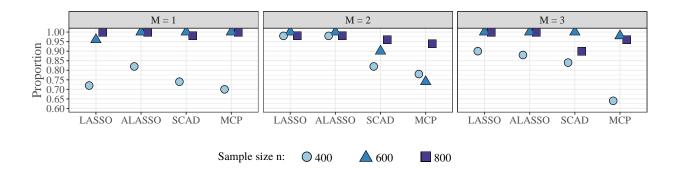


Figure 4: Proportion of correctly selected M using BIC based on 50 random samples for d = 20.

# 6 Case study: Identification of a recessionary regime

We consider a large-scale dataset of Canadian national and provincial macroeconomic indicators of observations from 1981 to 2022, containing 481 observations on 386 variables. The indicators belong to several categories, such as labour market, housing, construction and manufacturing. Fortin-Gagnon et al. (2018) analyzed a part of this dataset using probit and logit LASSO models to identify the turning points of three economic recession events, and predict recession probabilities. As MSVARs have been used to successfully model business cycles (Hamilton, 1988), our aim is to fit a model to this data and identify the recessionary regimes. The effects of manufacturing decline and house boom on US non-employment levels have been studied before (Charles et al., 2019). With this motivation, we consider housing price index (HPI) and manufacturing index (MI) for the top 5 Canadian provinces in terms of population<sup>1</sup>, hence the time series  $\mathbf{Y}_t$  has dimension d = 10. We fit MSVARs with M = 1, 2, 3, 4, p = 1, 2, 3, each penalty, and scenarios of sparse covariance and precision matrices. We use BIC and MDL, and the AR-order selection method based on the hierarchical group-LASSO, to choose the final model based on the ADALASSO with  $\widehat{M} = 3$ ,  $\widehat{p} = 1$  and sparse covariance matrices; see Supplement, Section A4 for all the details.

Figure 5 displays the estimated AR coefficients and covariance matrices for each regime, with transition probabilities  $(\hat{\alpha}_{11}, \hat{\alpha}_{12}) = (0.81, 0.13), (\hat{\alpha}_{21}, \hat{\alpha}_{22}) = (0.23, 0.69), \text{ and } (\hat{\alpha}_{31}, \hat{\alpha}_{32}) = (0.40, 0.40).$  We also show the estimated regime for each observation at time t as  $\arg \max_m \widehat{\mathbb{P}}(s_t = m | \mathbf{y}_{p+1:n})$ . In the same figure, we also mark the recession periods of the Canadian economy, which took place over the periods: June 1981 to October 1982, March 1990 to April 1992, and October 2008 to May 2009.

The logit model used in Fortin-Gagnon et al. (2018) provided smooth estimates of the predictive recession probabilities, which overlapped with the actual recessionary periods. From Figure 5, our fitted MSVAR is also able to identify a regime whose occurrence prob-

<sup>&</sup>lt;sup>1</sup>Statistics Canada, Table 17-10-0009-01; DOI: https://doi.org/10.25318/1710000901-eng

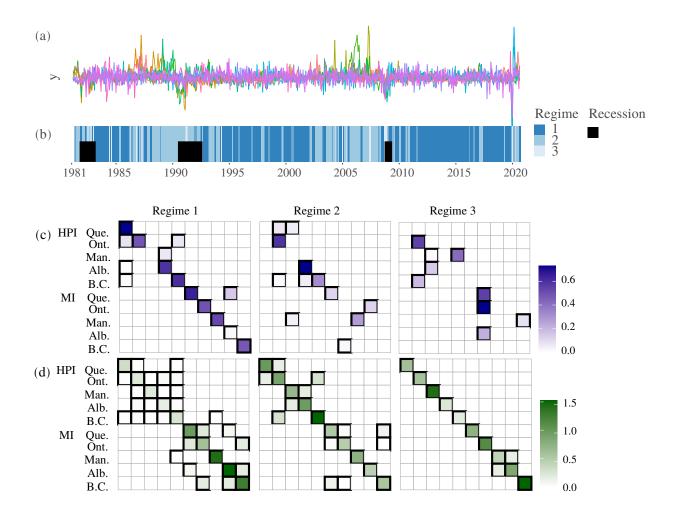


Figure 5: (a) Time series of d = 10 Canadian province macroeconomic indicators: 5 housing price (HPI) and 5 manufacturing (MI) indices. (b) Estimated most probable membership of each observation to any of  $\widehat{M} = 3$  regimes, and recession periods of Canadian economy. (c) Absolute value of estimated regime-AR coefficients, AR-order  $\widehat{p} = 1$ . (d) Absolute value of estimated regime-covariance matrix entries.

ability is substantially high before and during the recession periods. This corresponds to regime 2, and agrees with the general definition of a recession as a persistent decline in an economy with a specified minimum duration. The plot also shows that regime 3 captures particularly extreme values of some variables (HPI-Ont., HPI-Man., HPI-B.C., MI-Que., MI-Ont., MI-Alb.) and seems to appear more often around the recessionary periods. A proportion of 5% (25) observations are classified to regime 3. Figure 5(b) suggests that our method is able to detect the recession periods based on 2 indicators from 5 provinces, compared to 368 national and provincial variables used in Fortin-Gagnon et al. (2018).

**Prediction**. We consider the 8-month recessionary period over October 2008 to May 2009, and the subsequent non-recessionary 8-month period over June 2009 to January 2010, for a total of 16 test observations. We fit a sparse MSVAR model ( $\widehat{M} = 3$ ,  $\widehat{p} = 1$ ) using all the data up to the month prior to a test point under consideration, and perform an out-of-sample one-step-ahead regime-membership prediction. For a test point, we determine that it belongs to a recession if its predicted maximum regime probability corresponds to regimes either 2 or 3.

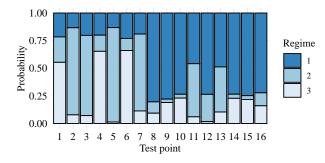


Figure 6: Out-of-sample one-step-ahead predicted regime membership probabilities for the Canadian province macroeconomic test data. The first 8 points correspond to the third Canadian recession, the last 8 points are non-recessionary.

Figure 6 shows the predicted probabilities for the 16 test observations. Note that even though we correctly identify the regimes for 14 test points, caution must be taken to predict recessions using the method since, as stated by Fortin-Gagnon et al. (2018), the presence of a recession is rather a rare event in the dataset.

# 7 Conclusion

We introduce a general penalized conditional likelihood method for fitting sparse MSVARs. It provides both sparse regime-specific AR coefficients and covariance or precision matrices, along with estimation of transition probabilities of the Markov chain. We prove consistency and sparsity properties of the proposed estimators. We also discuss estimation of the number of regimes and predictive density. We implement the method via an efficient modified EM algorithm. We examine the performance of the method via simulation and real data analysis. Simulations indicate that our method is reliable for fitting sparse MSVARs to moderate- and large-dimensional time series data. In practice, an increase in the number of regimes M and AR-order p has impact on the performance of the method, which is expected as the optimization problem (4) becomes more challenging. Our simulations also show that none of the penalties dominates the others in all the settings considered here. Thus, in practice, we suggest analyzing a dataset using all the penalties and choosing a sparse model that optimizes a selection criterion (e.g., BIC) as done in our real data analysis.

It can be shown that the BIC based on (18) does not under-estimate the true number of regimes M. A possible research direction is to investigate conditions under which BIC consistently estimates M. Another direction is the extension of our method to nonhomogeneous Markov chains. Shrinkage estimators can be incorporated by adjusting the penalty in (4) and also allowing the dimension d to increase with the sample size n as in the work of Fiecas et al. (2017), which require further investigation.

### Supplementary materials

A Supplementary pdf document to this paper includes: (i) additional methodology details including the EM algorithm, (ii) theoretical results and proofs, (iii) complementary numerical results, and (iv) complementary case study results.

Code: "code\_msvar.zip" contains the code to replicate Figure 1(a) and Figure 2(a).

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# Supplement to "Penalized estimation of sparse Markov regime-switching vector auto-regressive models"

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In this supplementary document, we provide complete details of the modified EM algorithm, proofs of the theoretical results on the asymptotic convergence and consistency of our penalized maximum likelihood estimates and predictive density estimates, and the results of additional numerical experiments not included in the main document. In the following, we denote the true model parameter vector as  $\boldsymbol{\theta}^{\star} \in \boldsymbol{\Theta} \subseteq \mathbb{R}^{K}$ , and denote a generic element of a parameter vector  $\boldsymbol{\theta}$  as  $\theta_{j}, j = 1, \ldots, K$ .

### A1 Methodology details

#### A1.1 Penalty functions

The well-studied choices of the penalty function  $r_{\lambda}$  in Section 3 of the main manuscript are:

- LASSO (L<sub>1</sub> norm):  $r_{\lambda}(\theta_j) = \lambda |\theta_j|;$
- Adaptive LASSO (ADALASSO, weighted L<sub>1</sub> norm):  $r_{\lambda}(\theta_j) = \lambda \widehat{w}_j |\theta_j|$ , with  $\widehat{w}_j = |\widehat{\theta}_j|^{-\gamma}$  and  $\gamma > 0$ , where  $\widehat{\theta}_j$  is a  $\sqrt{n}$ -consistent estimator of the true parameter  $\theta_j^{\star}$ ;
- SCAD:  $r_{\lambda}(\theta_j)$  is such that  $r'_{\lambda}(\theta) = \operatorname{sgn}(\theta) \left(\lambda \mathbb{1}_{\{|\theta| \leq \lambda\}} + \frac{(a\lambda |\theta|)_+}{(a-1)} \mathbb{1}_{\{\theta > \lambda\}}\right)$ , with parameter a > 2;
- MCP:  $r_{\lambda}(\theta_j)$  is such that  $r'_{\lambda}(\theta) = \operatorname{sgn}(\theta) \frac{(a\lambda |\theta|)_+}{a}$ , with parameter a > 1.

In Section A2 below we list general conditions on  $r_{\lambda}$  required to prove our theoretical results.

#### A1.2 Expectation (E-) step

Here we provide details of the forward-backward algorithm to compute the quantities

$$\zeta_{t,ij}^{(k)} = \mathbb{P}(\xi_{(t-1)i} = 1, \xi_{tj} = 1 | \mathbf{y}_{1:n}, \boldsymbol{\theta}^{(k)}, s_p), \quad \zeta_{ti}^{(k)} = \mathbb{P}(\xi_{ti} = 1 | \mathbf{y}_{1:n}, \boldsymbol{\theta}^{(k)}, s_p),$$

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at the (k+1)-th iteration of the EM algorithm. The forward recursion is also known as filtering, and the backward recursion as smoothing. We omit the terms  $(\boldsymbol{\theta}^{(k)}, s_p)$  from the notation throughout.

Recall the variable

$$oldsymbol{\xi}_t := egin{bmatrix} \xi_{t1} \ dots \ \xi_{t\mathrm{M}} \end{bmatrix} = egin{bmatrix} 1_{\{s_t=1\}} \ dots \ 1_{\{s_t=\mathrm{M}\}} \end{bmatrix}.$$

Let  $\iota_m$  denote the *m*-th column of the M-dimensional identity matrix. Using the model assumptions, we can write, for  $t \ge p + 1$ ,

$$\begin{split} \mathbb{P}(\mathbf{y}_t | \boldsymbol{\xi}_{t-1} = \boldsymbol{\iota}_i, \mathbf{y}_{1:t-1}) &:= \mathbb{P}(\mathbf{y}_t | \boldsymbol{\xi}_{(t-1)i} = 1, \mathbf{y}_{1:t-1}) \\ &= \sum_{m=1}^{M} \mathbb{P}(\mathbf{y}_t | \boldsymbol{\xi}_{(t-1)i} = 1, \boldsymbol{\xi}_{tm} = 1, \mathbf{y}_{1:t-1}) \mathbb{P}(\boldsymbol{\xi}_{tm} = 1 | \boldsymbol{\xi}_{(t-1)i} = 1, \mathbf{y}_{1:t-1}) \\ &= \sum_{m=1}^{M} \mathbb{P}(\mathbf{y}_t | \boldsymbol{\xi}_{tm} = 1, \mathbf{y}_{1:t-1}) \mathbb{P}(\boldsymbol{\xi}_{tm} = 1 | \boldsymbol{\xi}_{(t-1)i} = 1) \\ &= \sum_{m=1}^{M} \mathbb{P}(\mathbf{y}_t | \boldsymbol{\xi}_{tm} = 1, \mathbf{y}_{1:t-1}) \alpha_{im} = \sum_{m=1}^{M} \alpha_{im} \phi(\mathbf{y}_t; \boldsymbol{\mu}_t^{(m)}; \boldsymbol{\Sigma}^{(m)})). \end{split}$$

We collect the densities of  $\mathbf{y}_t$  conditional on  $\boldsymbol{\xi}_t$  and  $\mathbf{y}_{1:t-1}$  in the vector

$$\boldsymbol{\eta}_t := \begin{bmatrix} \mathbb{P}(\mathbf{y}_t | \boldsymbol{\xi}_t = \boldsymbol{\iota}_1, \mathbf{y}_{1:t-1}) \\ \vdots \\ \mathbb{P}(\mathbf{y}_t | \boldsymbol{\xi}_t = \boldsymbol{\iota}_{\mathrm{M}}, \mathbf{y}_{1:t-1}) \end{bmatrix} = \begin{bmatrix} \phi(\mathbf{y}_t; \boldsymbol{\mu}_t^{(1)}; \boldsymbol{\Sigma}^{(1)}) \\ \vdots \\ \phi(\mathbf{y}_t; \boldsymbol{\mu}_t^{(\mathrm{M})}; \boldsymbol{\Sigma}^{(\mathrm{M})}) \end{bmatrix}$$

Thus, we can rewrite the above conditional density as

$$\mathbb{P}(\mathbf{y}_t | \boldsymbol{\xi}_{t-1} = \boldsymbol{\iota}_i, \mathbf{y}_{1:t-1}) = \boldsymbol{\eta}_t^{\mathsf{T}} \mathbf{P}^{\mathsf{T}} \boldsymbol{\iota}_i.$$
(1)

As the regime-governing Markov chain is unobservable, the information at t-1 only consists of observations  $\mathbf{y}_{1:t-1}$  and not the regime indicator vector  $\boldsymbol{\xi}_{t-1}$ . For the purpose of estimation, we replace the vector  $\boldsymbol{\xi}_{t-1}$  by its conditional expected value, which in itself is estimated from the observed data as follows.

Denote the vectors

$$\widehat{\boldsymbol{\xi}}_{t|\tau} := \mathbb{E}(\boldsymbol{\xi}_t | \mathbf{y}_{1:\tau}) = \begin{bmatrix} \mathbb{P}(\boldsymbol{\xi}_t = \boldsymbol{\iota}_1 | \mathbf{y}_{1:\tau}) \\ \vdots \\ \mathbb{P}(\boldsymbol{\xi}_t = \boldsymbol{\iota}_{\mathrm{M}} | \mathbf{y}_{1:\tau}) \end{bmatrix} = \begin{bmatrix} \mathbb{P}(\xi_{t1} = 1 | \mathbf{y}_{1:\tau}) \\ \vdots \\ \mathbb{P}(\xi_{tM} = 1 | \mathbf{y}_{1:\tau}) \end{bmatrix}$$

for  $p \leq \tau \leq t$  and  $t \geq p + 1$ . Thus, using (1), the conditional probability density of  $\mathbf{y}_t$  given  $\mathbf{y}_{1:t-1}$  can be written as

$$egin{aligned} \mathbb{P}(\mathbf{y}_t|\mathbf{y}_{1:t-1}) &= \sum_{m=1}^{\mathrm{M}} \mathbb{P}(\mathbf{y}_t, oldsymbol{\xi}_{t-1} = oldsymbol{\iota}_m | \mathbf{y}_{1:t-1}) \ &= \sum_{m=1}^{\mathrm{M}} \mathbb{P}(\mathbf{y}_t|oldsymbol{\xi}_{t-1} = oldsymbol{\iota}_m, \mathbf{y}_{1:t-1}) \mathbb{P}(oldsymbol{\xi}_{t-1} = oldsymbol{\iota}_m | \mathbf{y}_{1:t-1}) \end{aligned}$$

$$= \sum_{m=1}^{M} \mathbb{P}(\mathbf{y}_t | \boldsymbol{\xi}_{(t-1)m} = 1, \mathbf{y}_{1:t-1}) \mathbb{P}(\boldsymbol{\xi}_{(t-1)m} = 1 | \mathbf{y}_{1:t-1})$$
$$= \boldsymbol{\eta}_t^{\mathsf{T}} \mathbf{P}^{\mathsf{T}} \sum_{m=1}^{M} \boldsymbol{\iota}_m \, \mathbb{P}(\boldsymbol{\xi}_{(t-1)m} = 1 | \mathbf{y}_{1:t-1}) = \boldsymbol{\eta}_t^{\mathsf{T}} \mathbf{P}^{\mathsf{T}} \widehat{\boldsymbol{\xi}}_{t-1|t-1}$$

On the other hand, for  $m = 1, \ldots, M$ , we have

$$\mathbb{P}(\xi_{tm} = 1 | \mathbf{y}_{1:t-1}) = \sum_{i=1}^{M} \mathbb{P}(\xi_{tm} = 1 | \xi_{(t-1)i} = 1, \mathbf{y}_{1:t-1}) \mathbb{P}(\xi_{(t-1)i} = 1 | \mathbf{y}_{1:t-1})$$
$$= \sum_{i=1}^{M} \alpha_{im} \mathbb{P}(\xi_{(t-1)i} = 1 | \mathbf{y}_{1:t-1}) = (\mathbf{P}^{\mathsf{T}} \widehat{\boldsymbol{\xi}}_{t-1|t-1})_{m},$$

where  $(\mathbf{v})_m$  refers to the *m*-th element of a vector  $\mathbf{v}$ . Thus, for all  $t \ge p + 1$ , we have

$$\widehat{\boldsymbol{\xi}}_{t|t-1} = \mathbf{P}^{\mathsf{T}} \widehat{\boldsymbol{\xi}}_{t-1|t-1},$$

and hence we get

$$\mathbb{P}(\mathbf{y}_t|\mathbf{y}_{1:t-1}) = oldsymbol{\eta}_t^\mathsf{T} \widehat{oldsymbol{\xi}}_{t|t-1} = \mathbf{1}_\mathrm{M}^\mathsf{T} \left(oldsymbol{\eta}_t \odot \widehat{oldsymbol{\xi}}_{t|t-1}
ight),$$

where  $\odot$  denotes the element-wise matrix multiplication and  $\mathbf{1}_{M} = (1, \dots, 1)^{\mathsf{T}}$ . Then

$$\widehat{\boldsymbol{\xi}}_{t|t} = \frac{\boldsymbol{\eta}_t \odot \widehat{\boldsymbol{\xi}}_{t|t-1}}{\mathbf{1}_{\mathrm{M}}^{\mathsf{T}} \left( \boldsymbol{\eta}_t \odot \widehat{\boldsymbol{\xi}}_{t|t-1} \right)},\tag{2}$$

which follows since, by Bayes' rule,

$$\mathbb{P}(\xi_{tm} = 1 | \mathbf{y}_{1:t}) = \mathbb{P}(\xi_{tm} = 1 | \mathbf{y}_t, \mathbf{y}_{1:t-1}) = \frac{\mathbb{P}(\mathbf{y}_t | \xi_{tm} = 1, \mathbf{y}_{1:t-1}) \mathbb{P}(\xi_{tm} = 1 | \mathbf{y}_{1:t-1})}{\mathbb{P}(\mathbf{y}_t | \mathbf{y}_{1:t-1})}$$

Now we consider  $\widehat{\pmb{\xi}}_{t|n}.$  First note that, by Bayes' rule and the model assumptions,

$$\begin{split} \mathbb{P}(\xi_{tm} = 1 | \xi_{(t+1)i} = 1, \mathbf{y}_{1:n}) &= \mathbb{P}(\xi_{tm} = 1 | \xi_{(t+1)i} = 1, \mathbf{y}_{1:t}, \mathbf{y}_{t+1:n}) \\ &= \frac{\mathbb{P}(\mathbf{y}_{t+1:n} | \xi_{tm} = 1, \xi_{(t+1)i} = 1, \mathbf{y}_{1:t}) \mathbb{P}(\xi_{tm} = 1 | \xi_{(t+1)i} = 1, \mathbf{y}_{1:t})}{\mathbb{P}(\mathbf{y}_{t+1:n} | \xi_{(t+1)i} = 1, \mathbf{y}_{1:t})} \\ &= \frac{\mathbb{P}(\mathbf{y}_{t+1:n} | \xi_{(t+1)i} = 1, \mathbf{y}_{1:t}) \mathbb{P}(\xi_{tm} = 1 | \xi_{(t+1)i} = 1, \mathbf{y}_{1:t})}{\mathbb{P}(\mathbf{y}_{t+1:n} | \xi_{(t+1)i} = 1, \mathbf{y}_{1:t})} \\ &= \mathbb{P}(\xi_{tm} = 1 | \xi_{(t+1)i} = 1, \mathbf{y}_{1:t}), \end{split}$$

which we use to write, once more using Bayes' rule,

$$\begin{split} \mathbb{P}(\xi_{tm} = 1, \xi_{(t+1)i} = 1 | \mathbf{y}_{1:n}) &= \mathbb{P}(\xi_{tm} = 1 | \xi_{(t+1)i} = 1, \mathbf{y}_{1:n}) \mathbb{P}(\xi_{(t+1)i} = 1 | \mathbf{y}_{1:n}) \\ &= \mathbb{P}(\xi_{tm} = 1 | \xi_{(t+1)i} = 1, \mathbf{y}_{1:t}) \mathbb{P}(\xi_{(t+1)i} = 1 | \mathbf{y}_{1:n}) \\ &= \frac{\mathbb{P}(\xi_{(t+1)i} = 1 | \xi_{tm} = 1, \mathbf{y}_{1:t}) \mathbb{P}(\xi_{tm} = 1 | \mathbf{y}_{1:t})}{\mathbb{P}(\xi_{(t+1)i} = 1 | \mathbf{y}_{1:t})} \mathbb{P}(\xi_{(t+1)i} = 1 | \mathbf{y}_{1:n}) \end{split}$$

$$=\frac{\mathbb{P}(\xi_{tm}=1|\mathbf{y}_{1:t})\mathbb{P}(\xi_{(t+1)i}=1|\xi_{tm}=1)}{\mathbb{P}(\xi_{(t+1)i}=1|\mathbf{y}_{1:t})}\mathbb{P}(\xi_{(t+1)i}=1|\mathbf{y}_{1:n}).$$
 (3)

Using the above expression, for m = 1, ..., M, we obtain

$$\begin{split} \mathbb{P}(\xi_{tm} = 1 | \mathbf{y}_{1:n}) &= \sum_{i=1}^{M} \mathbb{P}(\xi_{tm} = 1, \xi_{(t+1)i} = 1 | \mathbf{y}_{1:n}) \\ &= \sum_{i=1}^{M} \frac{\mathbb{P}(\xi_{tm} = 1 | \mathbf{y}_{1:t}) \mathbb{P}(\xi_{(t+1)i} = 1 | \xi_{tm} = 1)}{\mathbb{P}(\xi_{(t+1)i} = 1 | \mathbf{y}_{1:t})} \mathbb{P}(\xi_{(t+1)i} = 1 | \mathbf{y}_{1:n}) \\ &= \sum_{i=1}^{M} \frac{\alpha_{mi} \mathbb{P}(\xi_{(t+1)i} = 1 | \mathbf{y}_{1:n}) \mathbb{P}(\xi_{tm} = 1 | \mathbf{y}_{1:t})}{\mathbb{P}(\xi_{(t+1)i} = 1 | \mathbf{y}_{1:t})}. \end{split}$$

Collecting these into a vector, we get

$$\widehat{\boldsymbol{\xi}}_{t|n} = \begin{bmatrix} \mathbb{P}(\xi_{t1} = 1|\mathbf{y}_{1:n}) \\ \vdots \\ \mathbb{P}(\xi_{tM} = 1|\mathbf{y}_{1:n}) \end{bmatrix} = \begin{bmatrix} \mathbf{P}^{\mathsf{T}} \left( \widehat{\boldsymbol{\xi}}_{t+1|n} \oslash \widehat{\boldsymbol{\xi}}_{t+1|t} \right) \end{bmatrix} \odot \widehat{\boldsymbol{\xi}}_{t|t}, \tag{4}$$

denoting by  $\oslash$  the element-wise division.

From (2) and (4), using the current estimate  $\theta^{(k)}$ , we obtain the computations required in the E-step as follows:

• Forward recursion (filtering): for t = p + 1, ..., n,

$$\widehat{oldsymbol{\xi}}_{t|t} = rac{oldsymbol{\eta}_t \odot \widehat{oldsymbol{\xi}}_{t|t-1}}{oldsymbol{1}_{\mathrm{M}} \left(oldsymbol{\eta}_t \odot \widehat{oldsymbol{\xi}}_{t|t-1}
ight)} = rac{oldsymbol{\eta}_t \odot \mathbf{P}^{(k)} \widehat{oldsymbol{\xi}}_{t-1|t-1}}{oldsymbol{1}_{\mathrm{M}} \left(oldsymbol{\eta}_t \odot \mathbf{P}^{(k)} \widehat{oldsymbol{\xi}}_{t-1|t-1}
ight)},$$

where  $\odot$  denotes the element-wise product.

• Backward recursion (smoothing): for t = n - 1, ..., p + 1, after plugging in the filtered probabilities, we compute

$$\widehat{oldsymbol{\xi}}_{t|n} = \left[ \mathbf{P}^{(k)^{\mathsf{T}}} \left( \widehat{oldsymbol{\xi}}_{t+1|n} \oslash \widehat{oldsymbol{\xi}}_{t+1|t} 
ight) 
ight] \odot \widehat{oldsymbol{\xi}}_{t|t},$$

denoting by  $\oslash$  the element-wise division.

Finally, we set  $\zeta_{ti}^{(k)} = \mathbb{P}(\xi_{ti} = 1 | \mathbf{y}_{1:n})$  as the *i*-th entry of the vector  $\hat{\boldsymbol{\xi}}_{t|n}$ , for  $i = 1, \dots, M$ .

Regarding the joint probabilities  $\zeta_{t,ij}^{(k)} = \mathbb{P}(\xi_{(t-1)i} = 1, \xi_{tj} = 1 | \mathbf{y}_{1:n})$ , as discussed in (Krolzig, 1997, Chapter 5), they can be expressed in terms of the smoothed probabilities  $\hat{\xi}_{t|n,j}$  (*j*-th entry of  $\hat{\xi}_{t|l-1}$ ), the predicted probabilities  $\hat{\xi}_{t|l,j}$  (*j*-th entry of  $\hat{\xi}_{t|l-1}$ ), the filtered probabilities  $\hat{\xi}_{t-1|l-1,i}$  (*i*-th entry of  $\hat{\xi}_{t-1|l-1,i}$ ), and the transition probabilities estimates from the previous iteration. Using (3), the complete vector of joint probabilities, for  $t = p + 1, \ldots, n$ , is estimated by

$$\widehat{\boldsymbol{\xi}}_{t|n}^{\text{joint}} := \text{vec}(\mathbf{P}^{(k)}) \odot \left[ \left( \widehat{\boldsymbol{\xi}}_{t|n} \oslash \widehat{\boldsymbol{\xi}}_{t|t-1} \right) \otimes \widehat{\boldsymbol{\xi}}_{t-1|t-1} \right]$$

where  $\otimes$  denotes the Kronecker product, and whose (j-1)M + i entry corresponds to the term

$$\zeta_{t,ij}^{(k)} = \mathbb{P}(\xi_{(t-1)i} = 1, \xi_{tj} = 1 | \mathbf{y}_{1:n}) = rac{lpha_{ij}^{(k)}(\widehat{m{\xi}}_{t-1|t-1})_i(\widehat{m{\xi}}_{t|n})_j}{(\widehat{m{\xi}}_{t|t-1})_j},$$

for i, j = 1, ..., M. Recall  $(\mathbf{v})_i$  refers to the *i*-th element of a vector  $\mathbf{v}$ .

#### A1.3 Maximization (M-) step

#### A1.3.1 Transition probability matrix

The true transition probability matrix is estimated based on the joint distribution of  $\boldsymbol{\xi}_t, \boldsymbol{\xi}_{t+1}$  given the full sample  $\mathbf{y}_{1:n}$  as in (3). From (Krolzig, 1997, Equation 6.14), the estimate is given as

$$ext{vec}(\mathbf{P}^{(k+1)}) = \left[\widehat{oldsymbol{\xi}}^{ ext{joint}}
ight] \oslash \left[\mathbf{1}_{ ext{M}} \otimes \left(\left(\mathbf{1}_{ ext{M}}^{\mathsf{T}} \otimes \mathbf{I}_{ ext{M}}
ight) \widehat{oldsymbol{\xi}}^{ ext{joint}}
ight)
ight],$$

where  $\widehat{\boldsymbol{\xi}}^{\text{joint}} := \sum_{t=p+1}^{n} \widehat{\boldsymbol{\xi}}_{t|n}^{\text{joint}}$ , and  $\mathbf{I}_{\text{M}}$  is the M-dimensional identity matrix.

### A1.3.2 AR coefficient matrices

Recall the optimization problem with respect to the AR coefficients for regime m is given as

$$\min_{\{\mathbf{A}_{l}^{(m)}\}_{l=1}^{p}} \frac{1}{2(n-p)} \sum_{t=p+1}^{n} \zeta_{tm}^{(k)} (\mathbf{y}_{t} - \bar{\boldsymbol{\mu}}_{t}^{(m)})^{\mathsf{T}} \widehat{\boldsymbol{\Omega}}^{(m,k)} (\mathbf{y}_{t} - \bar{\boldsymbol{\mu}}_{t}^{(m)}) + \sum_{l=1}^{p} \sum_{i,j=1}^{d} r_{\lambda_{1}} (a_{l,ij}^{(m)}).$$

To solve it, we utilize the block-wise coordinate descent algorithm for VAR models suggested by Basu and Michailidis (2015). Let us first define

$$egin{aligned} \widehat{\Xi}_m &:= egin{bmatrix} \zeta_{(p+1)m}^{(k)} & \mathbf{0} \ & \ddots & \ & \mathbf{0} & & \zeta_{nm}^{(k)} \end{bmatrix}, \qquad \mathbf{y} &:= egin{bmatrix} (\mathbf{y}_{p+1} - oldsymbol{
u}^{(m,k+1)})^\mathsf{T} \ & dots \$$

Let  $\mathbf{y}_i^{(m)}$  and  $\mathbf{b}_i^{(m)}$  denote the *i*-th column and the *i*-th row of  $\widehat{\Xi}_m^{1/2} \mathbf{y}$  and  $[\mathbf{A}_1^{(m)}, \dots, \mathbf{A}_p^{(m)}]$ , respectively. Also let  $\mathbf{X}^{(m)} = \widehat{\Xi}_m^{1/2} \mathbf{X}$  and  $\widehat{\mathbf{\Omega}}^{(m,k)} = (\omega_{ij})_{1 \leq i,j \leq d}$ . The objective function can be rewritten as

$$\min_{\mathbf{b}_{i}^{(m)}, i=1,...,d} \frac{1}{2} \sum_{i=1}^{d} \sum_{j=1}^{d} \omega_{ij} \left( \mathbf{y}_{i}^{(m)} - \mathbf{X}^{(m)} \mathbf{b}_{i}^{(m)^{\mathsf{T}}} \right)^{\mathsf{T}} \left( \mathbf{y}_{j}^{(m)} - \mathbf{X}^{(m)} \mathbf{b}_{j}^{(m)^{\mathsf{T}}} \right) + \sum_{i=1}^{d} \sum_{j=1}^{dp} r_{\lambda_{1}} \left( b_{ij}^{(m)} \right),$$

where the  $b_{ij}^{(m)}$ , j = 1, ..., dp, are the elements of  $\mathbf{b}_i^{(m)}$ , i = 1, ..., d. As suggested in Basu and Michailidis (2015, Appendix C), we minimize the above objective function cyclically with respect to each  $\mathbf{b}_i^{(m)}$  until convergence. We repeat the following procedure for i = 1, ..., d, until convergence:

1. Set  $\mathbf{r}_{i} = \frac{1}{2\omega_{ii}} \sum_{j=1, j\neq i}^{d} \omega_{ij} \left( \mathbf{y}_{j}^{(m)} - \mathbf{X}^{(m)} \widehat{\mathbf{b}}_{j}^{\mathsf{T}} \right);$ 2. Update  $\widehat{\mathbf{b}}_{i} = \operatorname*{arg\,min}_{\mathbf{b}_{i}} \frac{\omega_{ii}}{2} \| \mathbf{y}_{i}^{(m)} + \mathbf{r}_{i} - \mathbf{X}^{(m)} \mathbf{b}_{i}^{\mathsf{T}} \|_{2}^{2} + \sum_{j=1}^{dp} r_{\lambda_{1}}(b_{ij}).$ 

To solve the optimization problem in Step 2 above, we employ a generalized gradient descent. In particular, we use a modified fast iterative-shrinkage thresholding algorithm (FISTA, Beck and Teboulle, 2009b), which was initially devised to solve optimization problems of the form  $\min_{\mathbf{b}} g(\mathbf{b}) + r_{\lambda}(\mathbf{b})$ . For a step size L, the iterative scheme is based on the update

$$\widehat{\mathbf{b}}_{i}^{(k+1)} = \operatorname*{arg\,min}_{\mathbf{b}_{i}} \left[ \frac{1}{2} \left\| \mathbf{b}_{i} - \left( \widehat{\mathbf{z}}_{i}^{(k)} - \mathrm{L}\omega_{ii} (\mathbf{y}_{i}^{(m)} + \mathbf{r}_{i} - \mathbf{X}^{(m)} (\widehat{\mathbf{z}}_{i}^{(k)})^{\mathsf{T}} \right)^{\mathsf{T}} \mathbf{X}^{(m)} \right) \right\|^{2} + \mathrm{L} \sum_{j=1}^{dp} r_{\lambda_{1}} (b_{ij}) \right]$$
(5)  
=:  $h(\widehat{\mathbf{z}}_{i}^{(k)}, \mathrm{L}),$ 

where  $\hat{\mathbf{z}}_{i}^{(k)}$  is an interpolation between  $\hat{\mathbf{b}}_{i}^{(k)}$  and  $\hat{\mathbf{b}}_{i}^{(k-1)}$ . Since the considered penalty function is decomposable with respect to individual elements of  $\mathbf{b}_{i}$ , the minimization problem above corresponds to a penalized least-squares problem with orthogonal predictors, and can be solved analytically for all the penalty functions we consider in this work. The exact formulas for this update for all the considered penalties are provided in Section A1.3.4.

Ideally, in an optimization algorithm the value of the objective function must not increase when computed over successive iterations. The original FISTA does not possess this property, making it vulnerable to divergence. This issue has been circumvented in another work by Beck and Teboulle (2009a), and we incorporate their approach in our implementation. Another enhancement we implement is known as FISTA with restart (Wen et al., 2017, and references therein), which resets the interpolation parameter every  $\kappa$  iterations for a pre-specified  $\kappa$ , and gives faster convergence compared to the original version of FISTA. See Algorithm 2 below.

We compute a step size L that ensures convergence as follows. First note that a Lipschitz constant of the smooth term in the objective function is  $\lambda_{\max}(\mathbf{X}^{\mathsf{T}}\widehat{\Xi}_m\mathbf{X})$ , where  $\lambda_{\max}(\mathbf{A})$  denotes the maximum eigenvalue of a real symmetric matrix  $\mathbf{A}$ . The knowledge of this constant provides a suitable step size (Beck and Teboulle, 2009b). For a given regime  $m \in \{1, \ldots, M\}$ , we can approximate the corresponding Lipschitz constant by observing the relation

$$\mathbf{X}^{\mathsf{T}}\widehat{\Xi}_m\mathbf{X} \preceq \mathbf{X}^{\mathsf{T}}\mathbf{X},$$

which holds since each element of the diagonal matrix  $\widehat{\Xi}_m$  lies in the interval (0, 1). Hence, following Guo et al. (2016, Theorem 4.4), we set L as any value satisfying

$$L < \begin{cases} \frac{2}{2C + \lambda_{\max}(\mathbf{X}^{\mathsf{T}}\mathbf{X})} & \text{if } C = 0, \\ \frac{2}{C + \lambda_{\max}(\mathbf{X}^{\mathsf{T}}\mathbf{X})} & \text{if } 0 < C \le \lambda_{\max}(\mathbf{X}^{\mathsf{T}}\mathbf{X})/2, \\ \frac{2}{2C + \sqrt{\lambda_{\max}(\mathbf{X}^{\mathsf{T}}\mathbf{X})^2 - C^2}} & \text{if } \lambda_{\max}(\mathbf{X}^{\mathsf{T}}\mathbf{X})/2 < C < \lambda_{\max}(\mathbf{X}^{\mathsf{T}}\mathbf{X}), \\ \frac{1}{C} & \text{if } \lambda_{\max}(\mathbf{X}^{\mathsf{T}}\mathbf{X}) = C, \end{cases}$$
(6)

where C is the weak-convexity constant of the penalty r, such that  $\sum_{j=1}^{dp} r_{\lambda_1}(b_{ij}) + \frac{C}{2} ||\mathbf{b}_i||^2$  is convex. This constant C exists for all the penalties considered in this work (see Section A1.3.4 below). By using such L we ensure the convergence of the algorithm to a local minimum. Our procedure is summarized in Algorithm 2.

#### A1.3.3 Covariance and precision matrices

Recall the optimization problem with respect to  $\mathbf{\Sigma}^{(m)}$  or  $\mathbf{\Omega}^{(m)}$  is

$$\min_{\boldsymbol{\Sigma}^{(m)} \succ 0} \frac{1}{2(n-p)} \Big( \widehat{n}_m \log |\boldsymbol{\Sigma}^{(m)}| + \operatorname{tr} \big( \boldsymbol{\Omega}^{(m)} \mathbf{S}^{(m)} \big) \Big) + \sum_{i \neq j=1}^d r_{\lambda_2} \big( \gamma_{ij}^{(m)} \big), \tag{7}$$

#### Algorithm 2 Monotone FISTA with restart for sparse VAR parameter estimation

$$\begin{array}{l} \text{Initialization: } \widehat{\mathbf{x}}_{i}^{1} = \widehat{\mathbf{b}}_{i}^{0}, \ t_{1} = 1, \ \kappa \geq 1, \ \text{L satisfying (6)}, \ \varepsilon > 0 \\ 1: \ \widehat{\mathbf{x}}_{i}^{k} = h(\widehat{\mathbf{z}}_{i}^{k}, \text{L}) \\ 2: \ \widehat{\mathbf{b}}_{i}^{k} = \arg\min\{h(\mathbf{x}, \text{L}) : \mathbf{x} \in \{\widehat{\mathbf{x}}_{i}^{k}, \widehat{\mathbf{b}}_{i}^{k-1}\} \} \\ 3: \ t_{k+1} = \frac{1 + \sqrt{1 + 4t_{k}^{2}}}{2} \\ 4: \ \widehat{\mathbf{z}}_{i}^{k+1} = \widehat{\mathbf{b}}_{i}^{k} + \left(\frac{t_{k-1}}{t_{k+1}}\right)(\widehat{\mathbf{b}}_{i}^{k} - \widehat{\mathbf{b}}_{i}^{k-1}) + \left(\frac{t_{k}}{t_{k+1}}\right)(\widehat{\mathbf{x}}_{i}^{k} - \widehat{\mathbf{b}}_{i}^{k}) \\ 5: \ \text{If } k \ \text{mod } \kappa = 0 \ \text{set } t_{k+1} = 1 \\ 6: \ \text{If } \|\widehat{\mathbf{b}}_{i}^{k} - \widehat{\mathbf{b}}_{i}^{k-1}\|/\|\widehat{\mathbf{b}}_{i}^{k-1}\| < \varepsilon \ \text{return } \widehat{\mathbf{b}}_{i}^{k}, \ \text{else } k = k+1 \ \text{and go to } 1 \end{array}$$

with  $\mathbf{\Gamma}^{(m)} = (\gamma_{ij}^{(m)})_{1 \leq i,j \leq d}$  being either  $\mathbf{\Sigma}^{(m)}$  or  $\mathbf{\Omega}^{(m)}$ . To simplify the notation, we drop the index *m* below. Loh and Wainwright (2015) proposed and theoretically studied an algorithm for regularized M-estimation for non-convex problems. Their algorithm performs a generalized gradient descent on the objective function, which is assumed to be the sum of a smooth function and a penalty function, similar to FISTA. We adopt their approach and implement it as follows. Let C be the weak-convexity constant of the penalty r, so that  $\sum_{i,j=1}^{d} r_{\lambda_2}(\mathbf{\Gamma}) + \frac{C}{2} \|\mathbf{\Gamma}\|_{\mathrm{F}}^2$  is convex (see Section A1.3.4). Next we provide the updates for the scenario where  $\mathbf{\Sigma}$  is penalized, followed by the updates in the scenario where the penalty is instead on  $\mathbf{\Omega}$ .

#### Penalization on the covariance matrix

The update with step size  $L_k$  for the optimization problem for penalized covariance is given as

$$\Sigma^{(k+1)} \in \underset{\Gamma \succ 0}{\operatorname{arg\,min}} \left\{ \frac{1}{2} \left\| \Gamma - \left( \Sigma^{(k)} - \mathcal{L}_k \left( \Omega^{(k)} - \Omega^{(k)} \mathbf{S} \Omega^{(k)} - \mathcal{C} \Sigma^{(k)} \right) \right) \right\|_{\mathcal{F}}^2 + \mathcal{L}_k \left( \sum_{i,j=1}^d r_{\lambda_2}(\gamma_{ij}) + \frac{\mathcal{C}}{2} \|\Gamma\|_{\mathcal{F}}^2 \right) \right\}$$

$$=: h_{\Sigma}(\Sigma^{(k)}, \mathcal{L}_k), \qquad (8)$$

which has a closed form for all the penalties we consider (see Section A1.3.4). Loh and Wainwright (2015) provided error bounds with respect to a global minimizer that hold with high probability, and stated that the iterates (8) quickly converge to a neighborhood of any global optimum under a set of smoothness and convexity conditions. Our objective function meets those requirements since its differentiable term has a Lipschitz-continuous gradient on a compact constraint set of the form  $\{\Sigma : \Sigma \succeq \delta I\}$  for some  $\delta > 0$ . This can be verified using ideas similar to those of Bien and Tibshirani (2011, Appendix 2). We compute the step size  $L_k$  using a backtracking line search; see, for example, Parikh and Boyd (2013, Section 4.2).

#### Penalization on the precision matrix

For the penalized precision matrix case, the difference arises on the differentiable part of the objective function and its gradient, in comparison with the covariance case. We perform each update as

$$\boldsymbol{\Omega}^{(k+1)} \in \underset{\boldsymbol{\Gamma} \succ 0}{\operatorname{arg\,min}} \left\{ \frac{1}{2} \left\| \boldsymbol{\Gamma} - \left( \boldsymbol{\Omega}^{(k)} - \mathcal{L}_{k} \left( -\boldsymbol{\Sigma}^{(k)} + \boldsymbol{S} - \mathcal{C}\boldsymbol{\Omega}^{(k)} \right) \right) \right\|_{\mathrm{F}}^{2} + \mathcal{L}_{k} \left( \sum_{i,j=1}^{d} r_{\lambda_{2}}(\gamma_{ij}) + \frac{\mathcal{C}}{2} \|\boldsymbol{\Gamma}\|_{\mathrm{F}}^{2} \right) \right\}$$

$$=: h_{\boldsymbol{\Omega}}(\boldsymbol{\Omega}^{(k)}, \mathcal{L}_{k}).$$
(9)

The theoretical properties of the iterative procedure as investigated by Loh and Wainwright (2015) are also applicable. The required Lipschitz continuity is implied when imposing the constraint  $\Sigma^{-1} \succeq \delta I$  for some specific  $\delta > 0$ . Such a  $\delta$  can be obtained with a derivation similar to Bien and Tibshirani (2011, Appendix 2). The final algorithm for solving (7) for a fixed regime m is outlined in Algorithm 3.

### Algorithm 3 Generalized gradient descent for sparse VAR covariance or precision matrix estimation

Initialization:  $\mathbf{\Sigma}^{(1)} = \mathbf{S}, \, \mathbf{\Omega}^{(1)} = \mathbf{S}^{-1}, \, \mathrm{L}_1 \leq 1, \, \varepsilon > 0$ 

- 1: Perform backtracking line search to obtain step size  $L_k$
- 2: (Penalized covariance)  $\Sigma^{(k+1)} = h_{\Sigma}(\Sigma^{(k)}, \mathbf{L}_{k})$ (Penalized precision)  $\Omega^{(k+1)} = h_{\Omega}(\Omega^{(k)}, \mathbf{L}_{k})$ 3: If  $\|\Sigma^{(k+1)} - \Sigma^{(k)}\|_{\mathbf{F}} / \|\Sigma^{(k)}\|_{\mathbf{F}} < \varepsilon$  or  $\|\Omega^{(k+1)} - \Omega^{(k)}\|_{\mathbf{F}} / \|\Omega^{(k)}\|_{\mathbf{F}} < \varepsilon$  return  $(\Sigma^{(k+1)}, \Omega^{(k+1)})$ else k = k + 1 and go to 1

#### A1.3.4 The proximal operators

The updates in Algorithm 2 and 3 of Section A1.3.3 have a closed-form formula for the penalty functions we consider, which we provide here. The update steps in (5), (8), and (9) can be obtained by solving the following

$$\min_x rac{1}{2}(x-z)^2 + 
u r_\lambda(x),$$

for  $x \in \mathbb{R} \setminus \{0\}$ , fixed  $z \in \mathbb{R}$  and  $\nu > 0$ . For example, in Algorithm 2, each element of  $\mathbf{b}_i$  as in (5), corresponds to x here. The formulas can be obtained by equating the subgradient of the objective function above to zero and solving for x. Thus, for a particular choice of the penalty function  $r_{\lambda}$ , the corresponding formulas for the updates in Algorithm 2 and 3 are given as:

$$\widehat{x}_{L_{1}} = \begin{cases} 0, & 0 \leq |z| \leq \nu\lambda, \\ z - \operatorname{sign}(z)\nu\lambda, & |z| \geq \nu\lambda; \end{cases}$$

$$\widehat{x}_{SCAD} = \begin{cases} 0, & 0 \leq |z| \leq \nu\lambda, \\ z - \operatorname{sign}(z)\nu\lambda, & \nu\lambda < |z| \leq (1+\nu)\lambda, \\ \frac{(a-1)z - \operatorname{sign}(z)a\nu\lambda}{a-1-\nu}, & (1+\nu)\lambda < |z| \leq a\lambda, \\ z, & |z| \geq a\lambda; \end{cases}$$

$$\widehat{x}_{MCP} = \begin{cases} 0, & 0 \leq |z| \leq \nu\lambda, \\ \frac{bz - \operatorname{sign}(z)b\nu\lambda}{b-\nu}, & \nu\lambda < |z| \leq b\lambda, \\ z, & |z| \geq b\lambda. \end{cases}$$

$$(10)$$

In the updates of Algorithm 2 and 3, we are required to set  $\nu = (\mathcal{L} + C)^{-1}$ , with C equal to 0,  $(a-1)^{-1}$  or  $b^{-1}$  for penalty functions L<sub>1</sub>-norm, SCAD or MCP, respectively; these are the weak convexity constants. For the adaptive LASSO, the update can be obtained by replacing  $\lambda$  in (10) with the weighted version  $\lambda \hat{w}$ , for a weight of the form  $\hat{w} = |\hat{\beta}|^{-\gamma}$ , where  $\hat{\beta}$  is a  $\sqrt{n}$ -consistent estimator of the parameter coordinate being estimated. We use the maximum-likelihood estimator with  $\gamma = 1$ . If the MLEs of the covariance or precision matrices are computationally singular, following the idea in ridge regression, we regularize them by adding a multiple of the identity matrix.

#### A1.4 Model initialization and identification

The performance of the modified EM algorithm is heavily reliant on an appropriate initialization. Our proposal here is motivated by the desired behaviour from a good estimate, of having a moderately high likelihood value. Define

$$\mathbf{y} := \begin{bmatrix} \mathbf{y}_{p+1}^{\mathsf{T}} \\ \vdots \\ \mathbf{y}_{n}^{\mathsf{T}} \end{bmatrix}, \qquad \mathbf{X} := \begin{bmatrix} \mathbf{X}_{p+1} \\ \vdots \\ \mathbf{X}_{n} \end{bmatrix}, \qquad \mathbf{X}_{t} := \begin{bmatrix} 1 \ \mathbf{y}_{t-1}^{\mathsf{T}} \ \mathbf{y}_{t-2}^{\mathsf{T}} \dots \mathbf{y}_{t-p}^{\mathsf{T}} \end{bmatrix}.$$

For  $m = 1, \ldots, M$ , we set the first iterates as

$$\begin{split} [\boldsymbol{\nu}^{(m,1)}, \mathbf{A}^{(m,1)}]^{\mathrm{T}} &= [\mathbf{X}^{\mathsf{T}} \mathbf{X}]^{-1} \mathbf{X}^{\mathsf{T}} \mathbf{y}, \\ \mathbf{\Sigma}^{(m,1)} &= \frac{1}{n-p} \left[ \mathbf{y} - \mathbf{X} [ \boldsymbol{\nu}^{(m,1)} \ \mathbf{A}^{(m,1)} ]^{\mathsf{T}} \right]^{\mathsf{T}} \left[ \mathbf{y} - \mathbf{X} [ \boldsymbol{\nu}^{(m,1)} \ \mathbf{A}^{(m,1)} ]^{\mathsf{T}} \right], \\ \mathbf{P}^{(1)} &= \alpha_{ij}^{(1)} = \begin{cases} \frac{1}{\mathrm{M}} + \delta, & i = j, \\ \frac{1}{\mathrm{M}-1} (1 - \alpha_{ii}^{(1)}), & i \neq j. \end{cases}$$

In other words, we use the least-squares and MLE estimates for a VAR(p) model (true p being assumed as given) as initial coefficient and covariance matrices for all the regimes. We use a generalized inverse if necessary. The transition probability matrix initial estimate is just a symmetric matrix, whose diagonal is specified by the quantity  $\delta \in (0, M^{-1})$ , and ensures that the estimates do not fall into a local maximum.

The interchange of regime labels in the MSVAR models during the estimation procedure can lead to a model identification problem. The model parameters belong to the same equivalence class if their regime labels are a permutation of any other label set in the class, and thus all the parameter combinations in an equivalence class lead to the same model. After convergence of the EM algorithm, we perform a permutation of the regime labels, as suggested by Krolzig (1997), according to a prespecified ordering on the magnitude of the estimated VAR coefficients. For example, different regimes labels can be indexed based on the increasing order of mean of the intercept vector. This ensures that the labels of the final estimates always follow the pre-specified order which takes care of the model identifiability. Another requirement of model identification is the existence of a well-defined distribution function. This condition is trivially satisfied as we assume Gaussian noise in our MSVAR models.

### A1.5 Selection of the penalty parameters $\lambda_1$ and $\lambda_2$

Inspired by Zhang et al. (2010), we select the penalty parameters by optimizing a model selection criterion, which is composed of a loss function and a model complexity term which involves the number of degrees of freedom of a candidate model. Here, we assume a fixed number of regimes M and AR-order p.

Let  $\widehat{\theta}_n(\lambda)$  be the MPLE given a pair  $\lambda = (\lambda_1, \lambda_2) \in \mathcal{I} \times \mathcal{I} \subset \mathbb{R}^2$ , for a pre-specified set of values  $\mathcal{I}$ . We denote the regime-specific degrees of freedom of the corresponding fitted MSVAR model as

$$D_m(\widehat{\theta}_n(\lambda)) = \sum_{k=1}^p \sum_{i,j=1}^d \mathbb{1}_{\{(\widehat{a}_{k,ij}^{(m)}) \neq 0\}} + \sum_{i,j=1}^d \mathbb{1}_{\{\widehat{\gamma}_{ij}^{(m)} \neq 0\}}, \quad m = 1, \dots, M,$$

and define the selection criterion

$$\mathcal{C}_1(oldsymbol{\lambda}) = -2l_n(\widehat{oldsymbol{ heta}}_n(oldsymbol{\lambda}); s_p) + c\sum_{m=1}^{\mathrm{M}} \mathrm{D}_m(\widehat{oldsymbol{ heta}}(oldsymbol{\lambda})),$$

where  $l_n$  is the conditional log-likelihood and c is a constant controlling the model complexity. A similar criterion, denoted by C, is used in the main manuscript for the selection of the number of regimes M. In the above criterion, if c = 2 or  $c = \log(n-p)$ , we obtain the AIC or BIC, respectively. A data-dependent choice of the tuning parameters  $\lambda = (\lambda_1, \lambda_2)$  is

$$\boldsymbol{\lambda} \in \underset{\boldsymbol{\lambda}_0 \in \mathcal{I} \times \mathcal{I}}{\operatorname{arg\,min}} \mathcal{C}_1(\boldsymbol{\lambda}_0). \tag{11}$$

The usual practice is to take  $\mathcal{I}$  to be a discretization of the set  $[0, \tilde{\lambda}]$  for some  $\tilde{\lambda} > 0$ , and choose  $\boldsymbol{\lambda} = (\lambda_1, \lambda_2)$  as in (11). In our numerical studies, the pairs  $(\lambda_1, \lambda_2)$  take values on a grid with increments of 0.25 on the logarithmic scale.

We propose a computationally efficient coordinate-descent approach which optimizes  $C_1$  by performing a cyclical search on either coordinate of the argument  $\lambda_0 \in \mathcal{I} \times \mathcal{I}$ , while keeping the other fixed until reaching convergence. To increase the accuracy, we perform additional searches at different resolutions of  $\mathcal{I}$ . Thus, the coarsest grid corresponds to a global coordinate descent, whereas subsequent searches are performed at decreasing resolutions in a neighborhood of the optimum grid point found in the previous resolution. This refines the result further. The discrete nature of the problem enables this procedure to converge to a local optimum. For a step size s > 0, window radius  $w \in \mathbb{N}$ , and center  $\lambda > 0$ , let  $\mathcal{I}_w(s, \lambda) := \{\max\{\lambda - ws, 0\}, \ldots, \lambda - 2s, \lambda - s, \lambda, \lambda + s, \lambda + 2s, \ldots, \min\{\lambda + ws, \tilde{\lambda}\}\}$ . Algorithm 4 summarizes the selection procedure of the penalty tuning parameters and number of regimes.

Algorithm 4 Selection of Tuning Parameters and Number of Regimes

1: Input:  $M_{\max} \in \mathbb{N}, \, \widetilde{\lambda} > 0, \, \widehat{\lambda}_2^0 = 0$ 2: For  $M = 1, \ldots, M_{max}$  do For k = 1, 2, ... do 3:  $\begin{aligned} \widehat{\lambda}_{1}^{k} &\in \underset{\lambda_{1} \in \mathcal{I}_{\infty}(s, \widetilde{\lambda}/2)}{\arg\min} \mathcal{C}_{1}(\lambda_{1}, \widehat{\lambda}_{2}^{k-1}), \\ \widehat{\lambda}_{2}^{k} &\in \underset{\lambda_{2} \in \mathcal{I}_{\infty}(s, \widetilde{\lambda}/2)}{\arg\min} \mathcal{C}_{1}(\widehat{\lambda}_{1}^{k}, \lambda_{2}), \end{aligned}$ until  $\widehat{\lambda}_{i}^{k} = \widehat{\lambda}_{i}^{k-1}, \ i = 1, 2$ Set  $c_i = \widehat{\lambda}_i^k, i = 1, 2$ For j = 1, 2 do  $\widehat{\lambda}_2^0 = c_2$ For  $k = 1, 2, \dots$  do 4: 5: 6: 7:  $\begin{aligned} \widehat{\lambda}_{1}^{k} &\in \underset{\lambda_{1} \in \mathcal{I}_{w}(s/2^{j},c_{1})}{\arg\min} \mathcal{C}_{1}(\lambda_{1},\widehat{\lambda}_{2}^{k-1}), \\ \widehat{\lambda}_{2}^{k} &\in \underset{\lambda_{2} \in \mathcal{I}_{w}(s/2^{j},c_{2})}{\arg\min} \mathcal{C}_{1}(\widehat{\lambda}_{1}^{k},\lambda_{2}), \end{aligned}$  $ext{until } \widehat{\lambda}_i^k = \widehat{\lambda}_i^{k-1}, \ i = 1,2 \ ext{Set } c_i = \widehat{\lambda}_i^k, \ i = 1,2 \ ext{Set } c_i = \widehat{\lambda}_i^k, \ i = 1,2 \ ext{}$ 8: Set  $\boldsymbol{\lambda}_{\mathrm{M}} = (\widehat{\lambda}_{1}^{k}, \widehat{\lambda}_{2}^{k})$ 9: Compute  $\mathcal{C}(M)$  defined in the main manuscript using  $\boldsymbol{\lambda}_{M}$ 10: 11: Set  $\mathbf{M} = \arg\min\{\mathcal{C}(1), \ldots, \mathcal{C}(\mathbf{M}_{\max})\}$ 12: Return  $(\boldsymbol{\lambda}_{\widehat{M}}, \widehat{M})$ 

### A1.6 AR-order selection

By adapting the approach for VARs by Nicholson et al. (2020), we devise a procedure for AR-order selection that has a superior performance for our models compared to a criterion-based approach. It is based on penalizing the AR coefficients using the group LASSO so that the spurious lag entries are shrunk to zero. This AR-order selection procedure, described next, is straightforward to incorporate into Algorithm 4.

First note that the AR-order selection can be done by relying on the consistency of the penalized estimator. We can consider a pre-specified AR-order  $p_0 \ge p$ , and express an MSVAR model with AR-order  $p_0$  as an MSVAR model with AR-order  $p_0$ , with  $\mathbf{A}_i^{(m)} \equiv 0_{d \times d}$  for i > p and  $m = 1, \ldots, M$ . However, a methodologically more suitable approach is to consider the group LASSO (Yuan and Lin, 2006) as done for VAR models by Song and Bickel (2011) and Nicholson et al. (2020).

Group LASSO shrinks all the elements in a given group simultaneously towards zero, that is, all the coefficients in a group are simultaneously estimated as either zero or nonzero. Nicholson et al. (2020) proposed a series of hierarchical-lag penalties derived from the group LASSO. The groups are designed in a nested manner so that if the group corresponding to lag i is set to zero, then its nested groups, corresponding to a lag greater than i, remain zero. Among the various hierarchical penalties, the element-wise hierarchical penalty is the most suitable for our models as we do not assume any particular sparsity structure on the AR coefficient matrices. It is also straightforward to optimize in our context. For regime m, we write

$$\min_{\{\mathbf{A}_{l}^{(m)}\}_{l=1}^{p}} \frac{1}{2(n-p)} \sum_{t=p+1}^{n} \zeta_{tm}^{(k)} (\mathbf{y}_{t} - \bar{\boldsymbol{\mu}}_{t}^{(m)})^{\mathsf{T}} \widehat{\boldsymbol{\Omega}}^{(m,k)} (\mathbf{y}_{t} - \bar{\boldsymbol{\mu}}_{t}^{(m)}) + \mathrm{R}(\mathbf{A}_{1}^{(m)}, \dots, \mathbf{A}_{p_{0}}^{(m)}; \lambda), \\ \mathrm{R}(\mathbf{A}_{1}^{(m)}, \dots, \mathbf{A}_{p_{0}}^{(m)}; \lambda) = \lambda \sum_{j=1}^{d} \sum_{k=1}^{d} \sum_{i=1}^{p} \sqrt{p_{0} - i + 1} \left( \sum_{\ell=i}^{p_{0}} (\mathbf{A}_{\ell}^{(m)})_{jk}^{2} \right)^{1/2},$$

so that each group indexed by lag *i* contains the element (j, k) of all matrices of lag  $\ell \ge i$ . Furthermore, the adaptive version of the hierarchical penalty is straightforward to obtain as an extension of the adaptive group LASSO (Wang and Leng, 2008). After estimation we can set

$$\widehat{p} = \max\{k \in \{1, \dots, p_0\} : \widehat{\mathbf{A}}_k^{(m)} \neq \mathbf{0}_{d \times d}\}.$$
(12)

An advantage of this AR-order selection method is that the penalty, and therefore the optimization problem, can be decomposed along coordinates, which enables us to apply the optimization framework we employ for the penalized AR coefficients estimation in Section A1.3.2.

We examine the performance of the above AR-order selection method via simulations. We set M = 2, d = 20, and sample sizes  $n \in \{400, 600, 800\}$ . We consider the true AR-orders p = 1, 2, 3, and set the over-specified candidate  $p_0 = 4$ . We compute the mean of the performance measure  $\mathbb{1}_{\{\widehat{p}=p\}}$ , with  $\widehat{p}$  obtained using (12). For both sparsity scenarios **S1** (sparse covariance matrices) and **S2** (sparse precision matrices), we obtain a mean performance measure of the true AR-order selection of at least 0.90. It is worth mentioning that using a AR-order selection procedure based on the BIC did not provide satisfactory results.

## A2 Theoretical results

In this section, we study the large sample properties of the MPLE  $\hat{\theta}_n$  introduced in Section 3 of the main manuscript, when the true number of regimes M is correctly specified or over-estimated. We denote the true MSVAR model parameter vector by  $\theta^*$ , and partition it as  $\theta^* = [(\theta^{*A})^{\mathsf{T}}, (\theta^{*A^c})^{\mathsf{T}}]^{\mathsf{T}}$ , so that  $\theta^{*A^c} = \mathbf{0}$ , and  $\theta^{*A}$  is a subvector of all the remaining non-zero parameters. Here,  $\mathcal{A}$  also denotes the set of indices k of all active (nonzero) parameters, and its complement  $\mathcal{A}^c$ , similarly contains the indices of all inactive (zero) parameters. Then, the cardinality of the index set of a parameter vector  $\theta$ , denoted as  $|\mathcal{A} \cup \mathcal{A}^c|$ , is equal to  $\mathsf{K} = \mathsf{M}(d + pd^2 + d(d + 1)/2) + \mathsf{M}(\mathsf{M} - 1)$ . Furthermore, we let  $\mathcal{E} \subset \mathcal{A} \cup \mathcal{A}^c$ , denote the set of indices k of the model parameters on which we perform the penalization. Recall, that we do not penalize the parameters corresponding to the transition probability matrix  $\mathbf{P}$  of the hidden Markov chain, the AR intercepts  $\boldsymbol{\nu}^{(m)}$ , and the diagonal entries of the covariance matrices  $\boldsymbol{\Sigma}^{(m)}, m = 1, \ldots, \mathsf{M}$ . To establish our theoretical results, we require certain conditions on the process  $\{\mathbf{Y}_t\}$ , the penalty function  $r_{\lambda_n}$  used in (4) of the main manuscript, and the tuning parameter  $\lambda_n$ . Let  $r'_{\lambda_n}(\cdot)$  and  $r''_{\lambda_n}(\cdot)$  denote the first and second derivatives, respectively, of the function  $r_{\lambda_n}(\cdot)$  with respect to  $\theta$ .

Assumption 1. For all  $\lambda \in \mathbb{R}$ ,  $r_{\lambda}(0) = 0$ . The function  $r_{\lambda}(\theta)$  is continuous, symmetric, nonnegative, nondecreasing, continuously differentiable for all  $|\theta| > 0$ , and twice continuously differentiable for all  $|\theta| > c\lambda$  for some constant c > 0.

Assumption 2. As  $n \to \infty$ ,  $\lambda_n = o(1)$ , and for true value of model parameters  $\theta_k^{\star}$ ,

$$a_n = \max_{k \in \mathcal{A} \cap \mathcal{E}} \{ |r'_{\lambda_n}(\theta_k^{\star})| \} = \mathcal{O}(n^{-1/2}), \qquad \qquad b_n = \max_{k \in \mathcal{A} \cap \mathcal{E}} \{ |r''_{\lambda_n}(\theta_k^{\star})| \} = o(1)$$

Assumption 3.  $\liminf_{n \to \infty} \liminf_{|\theta| = \mathrm{O}(n^{-1/2})} \sqrt{n} r'_{\lambda_n}(\theta) = +\infty.$ 

Assumption 1 imposes a differentiability condition on the penalty function which allows to perform the Taylor expansion of the objective function around the true optimizer. Assumption 2 is essential to show the existence of a  $\sqrt{n}$ -consistent optimizer of the penalized conditional loglikelihood function, and Assumption 3 is required to prove the sparsity property for the estimator of the model parameter. For simplicity of exposition, we suppose that the penalty function  $r_{\lambda}$  used for the covariance (or precision) matrix parameters is the same as for the VAR matrix coefficients, but this is not a requirement. In addition, in contrast to the related work of Monbet and Ailliot (2017), we employ a unique tuning parameter  $\lambda_n$  for all the regimes; the computational burden is alleviated in this way and our experiments show that one tuning parameter for all the regimes also performs well. As we consider the situation that for any  $\boldsymbol{\theta} \in \boldsymbol{\Theta}$ , we have  $\max_{m \in \{1,...,M\}} \lambda_{\min} (\Sigma_m) \geq \delta > 0$ , we do not penalize for a possibly singular  $\Sigma_m$ . Here,  $\lambda_{\min}(\mathbf{A})$  denotes the smallest eigenvalue of a real symmetric matrix  $\mathbf{A}$ . We assume that the true number of regimes in the MSVAR model is already given.

We use the results on consistency and asymptotic normality of the maximum-likelihood estimators obtained by Douc et al. (2004) for the MSVAR models. All their assumptions follow in our setting of a homogeneous regime-governing Markov chain  $S_t$  and the Gaussian noise process.

**Proposition 6.1.** (Douc et al., 2004, Proposition 1-3) There exists a deterministic function  $l(\cdot)$  such that

(i) for all  $s_p \in \{1, \ldots, M\}$ , and  $\boldsymbol{\theta} \in \boldsymbol{\Theta}$ ,

$$\lim_{n\to\infty} n^{-1} l_n(\boldsymbol{\theta}; s_p) = l(\boldsymbol{\theta}), \quad \mathbb{P}_{\boldsymbol{\theta}^{\star}}\text{-a.s. and in } \mathrm{L}^1(\mathbb{P}_{\boldsymbol{\theta}^{\star}});$$

- (ii)  $\lim_{n\to\infty} \sup_{\theta\in\Theta} \max_{s_p} |n^{-1}l_n(\theta; s_p) l(\theta)| = 0$ ,  $\mathbb{P}_{\theta^{\star}}$ -a.s.;
- (iii)  $l(\boldsymbol{\theta}) \leq l(\boldsymbol{\theta}^{\star})$  and  $l(\boldsymbol{\theta}) = l(\boldsymbol{\theta}^{\star})$  if and only if  $\boldsymbol{\theta} = \boldsymbol{\theta}^{\star}$ .

The above results ensure the strong consistency of the conditional MLE estimator. Next, we also have a central limit theorem for the score function and a law of large numbers for the observed Fisher information.

#### Proposition 6.2.

(i) (Douc et al., 2004, Theorem 2) (Central limit theorem) For any  $s_p \in \{1, \dots, M\}$ 

$$n^{-1/2} \nabla_{\boldsymbol{\theta}} l_n(\boldsymbol{\theta}^{\star}; s_p) \stackrel{\mathbb{P}_{\boldsymbol{\theta}^{\star}}}{\longrightarrow} \mathcal{N}(\mathbf{0}, \mathrm{I}(\boldsymbol{\theta}^{\star})),$$

where  $I(\theta^*)$  is the asymptotic Fisher information matrix, defined as the covariance of asymptotic score function.

(ii) (Douc et al., 2004, Theorem 3) (Law of large numbers) For any  $s_p \in \{1, \ldots, M\}$  and a possibly stochastic sequence  $\{\boldsymbol{\theta}_n^{\star}\} \in \boldsymbol{\Theta}$  converging to  $\boldsymbol{\theta}^{\star} \mathbb{P}_{\boldsymbol{\theta}\star}$ -almost surely, we have

$$-n^{-1}\nabla^2_{\boldsymbol{\theta}} l_n(\boldsymbol{\theta}_n^\star;s_p) \longrightarrow \mathrm{I}(\boldsymbol{\theta}^\star), \mathbb{P}_{\boldsymbol{\theta}\star}\text{-a.s.}$$

As fixing the initial state of the regime-governing Markov chain does not affect our asymptotic convergence results, we simplify our notation for  $l_n(\boldsymbol{\theta}; s_p)$  and  $\mathcal{L}_n(\boldsymbol{\theta}; s_p)$  by rewriting them as  $l_n(\boldsymbol{\theta})$ and  $\mathcal{L}_n(\boldsymbol{\theta})$ , respectively. The following results hold for any  $s_p \in \{1, \ldots, M\}$ . We first show that under the correct specification of M and appropriate choices of  $(r_{\lambda_n}, \lambda_n)$ , there exists a local maximizer  $\hat{\boldsymbol{\theta}}_n$  of the penalized conditional likelihood function  $\mathcal{L}_n(\boldsymbol{\theta})$  that is a consistent and sparse estimator of  $\boldsymbol{\theta}^{\star}$ .

**Theorem 1.** Suppose that  $\{\mathbf{Y}_t\}$  is generated according to a stationary and ergodic MSVAR process. Further assume that Assumption 1 - 3 on the tuning parameter  $\lambda_n$  and penalty function  $r_{\lambda_n}$  hold. Then, as  $n \to \infty$ ,

(i) there exists a local maximizer  $\widehat{\boldsymbol{\theta}}_n$  of the penalized log-likelihood  $\mathcal{L}_n(\boldsymbol{\theta})$  such that  $\|\widehat{\boldsymbol{\theta}}_n - \boldsymbol{\theta}^\star\| = O_P(n^{-1/2});$ (ii) for any  $\sqrt{n}$ -consistent estimator  $\widehat{\boldsymbol{\theta}}_n$  of the true sparse parameter  $\boldsymbol{\theta}^\star$  with  $\boldsymbol{\theta}^{\star \mathcal{A}^c} = \mathbf{0}$ , we have  $\mathbb{P}(\widehat{\boldsymbol{\theta}}_n^{\mathcal{A}^c} = \mathbf{0}) \to 1.$ 

*Proof.* Let  $\alpha_n = n^{-1/2} + a_n$ . For the first claim, it suffices to show that for  $n \ge n_0$  with  $n_0$  large enough, for any  $\varepsilon > 0$  there exists a large constant C such that

$$\mathbb{P}\Big(\sup_{\|\mathbf{u}\|=\mathcal{C}} \mathcal{L}_n(\boldsymbol{\theta}^{\star} + \alpha_n \mathbf{u}) < \mathcal{L}_n(\boldsymbol{\theta}^{\star})\Big) \ge 1 - \varepsilon.$$
(13)

In other words, we need to show that for  $\|\mathbf{u}\| = C$ , as  $n \to \infty$ ,  $\mathcal{L}_n(\boldsymbol{\theta}^* + \alpha_n \mathbf{u}) - \mathcal{L}_n(\boldsymbol{\theta}^*) < 0$ uniformly in  $\mathbf{u}$  with probability approaching to one. As the penalty function is non-negative, for  $k \in \mathcal{A}^c$ ,  $r_{\lambda_n}(\boldsymbol{\theta}_k^* + \alpha_n u_k) \geq 0$ . Also recall that  $r_{\lambda_n}(0) = 0$ . Then, we have the following inequality:

$$\begin{split} \mathrm{V}_{n}(\mathbf{u}) &\equiv \left(\mathcal{L}_{n}(\boldsymbol{\theta}^{\star} + \alpha_{n}\mathbf{u}) - \mathcal{L}_{n}(\boldsymbol{\theta}^{\star})\right) \\ &= \frac{1}{(n-p)}l_{n}(\boldsymbol{\theta}^{\star} + \alpha_{n}\mathbf{u}) - \sum_{k\in\mathcal{E}}r_{\lambda_{n}}\left(\theta_{k}^{\star} + \alpha_{n}u_{k}\right) - \frac{1}{(n-p)}l_{n}(\boldsymbol{\theta}^{\star}) + \sum_{k\in\mathcal{E}}r_{\lambda_{n}}\left(\theta_{k}^{\star}\right) \\ &\leq \frac{1}{(n-p)}\left(l_{n}(\boldsymbol{\theta}^{\star} + \alpha_{n}\mathbf{u}) - l_{n}(\boldsymbol{\theta}^{\star})\right) - \sum_{k\in\mathcal{A}\cap\mathcal{E}}\left(r_{\lambda_{n}}\left(\theta_{k}^{\star} + \alpha_{n}u_{k}\right) - r_{\lambda_{n}}\left(\theta_{k}^{\star}\right)\right) \\ &=:\mathrm{D}_{1n}(\mathbf{u}) - \mathrm{D}_{2n}(\mathbf{u},\lambda_{n}). \end{split}$$

We use the Taylor expansion of the likelihood function  $l_n$  to write:

$$\mathrm{D}_{1n}(\mathbf{u}) = rac{lpha_n}{(n-p)} \mathbf{u}^\mathsf{T} 
abla_ heta l_n(oldsymbol{ heta}^\star) + rac{1}{2(n-p)} lpha_n^2 \mathbf{u}^\mathsf{T} 
abla_ heta^2 l_n(\widetilde{oldsymbol{ heta}}_n) \mathbf{u}.$$

In the above  $\tilde{\boldsymbol{\theta}}_n$  is equal to  $s\boldsymbol{\theta}^{\star} + (1-s)(\boldsymbol{\theta}^{\star} + \alpha_n \mathbf{u})$  for some 0 < s < 1. From Proposition 6.2(i) we have for large enough n,  $\nabla_{\theta} l_n(\boldsymbol{\theta}^{\star}) = O_P(n^{1/2})$ . This gives  $\alpha_n \nabla_{\theta} l_n(\boldsymbol{\theta}^{\star}) = O_P(\alpha_n n^{1/2}) = O_P(\alpha_n^2 n)$ , due to Assumption 2 on  $a_n$ . From Proposition 6.2(ii), for large enough n, we have  $-\frac{1}{n} \nabla_{\theta}^2 l_n(\boldsymbol{\theta}) = I(\boldsymbol{\theta}^{\star})(1 + o_P(1))$ . Then, we get

$$D_{1n}(\mathbf{u}) = \frac{1}{(n-p)} \|\mathbf{u}\| O_{\mathrm{P}}(\alpha_n^2 n) - \frac{1}{2(n-p)} \mathbf{u}^{\mathsf{T}} \mathrm{I}(\boldsymbol{\theta}^{\star}) \mathbf{u} n \alpha_n^2 (1+o_{\mathrm{P}}(1)).$$
(14)

In the above, it is clear that the second term dominates the first uniformly for  $\|\mathbf{u}\| = C$  with a sufficiently large C. Recall that for all  $k \in \mathcal{A}, \theta_k^* \neq 0$  and for large enough  $n, |\theta_k^*| > c\lambda_n$  for some constant c > 0, since  $\lambda_n = o(1)$  due to Assumption 2. As  $\alpha_n = o(1)$ , we can perform a second-order Taylor expansion in  $D_{2n}(\mathbf{u}, \lambda_n)$  due to Assumption 1, and obtain the following inequality,

$$D_{2n}(\mathbf{u},\lambda_n) = \sum_{k \in \mathcal{A} \cap \mathcal{E}} \left( \alpha_n r'_{\lambda_n} \left( \theta_k^{\star} \right) u_k + \frac{\alpha_n^2}{2} r''_{\lambda_n} \left( \theta_k^{\star} \right) u_k^2 \left( 1 + o(1) \right) \right)$$

$$\leq \sqrt{|\mathcal{A} \cup \mathcal{A}^c|} \alpha_n a_n \|\mathbf{u}\| + \frac{\alpha_n^2}{2} b_n \|\mathbf{u}\|^2.$$
(15)

From the assumption on the rates of decay of  $a_n$  and  $b_n$  in Assumption 2, the above upper bound can also be dominated by the second term of (14) by choosing a large enough C. Thus, from the results in (14) and (15), we can conclude that for sufficiently large n,  $V_n(\mathbf{u}) < 0$ , and that the hypothesis in (13) holds.

To prove the result on the oracle property of the penalized estimator in Theorem 1(ii), we first show the following lemma.

**Lemma 6.3.** Suppose that  $\boldsymbol{\theta}_n = [(\boldsymbol{\theta}_n^{\mathcal{A}})^{\mathsf{T}}, (\boldsymbol{\theta}_n^{\mathcal{A}^c})^{\mathsf{T}}]^{\mathsf{T}}$ , is such that  $\|\boldsymbol{\theta}_n - \boldsymbol{\theta}^{\star}\| = O_{\mathsf{P}}(n^{-1/2})$  as  $n \to \infty$ . Then, for tuning parameter  $\lambda_n$  and penalty function  $r_{\lambda_n}$  satisfying Assumption 1–3, we have as  $n \to \infty$ , with probability tending to one that

$$\boldsymbol{\mathcal{L}}_n\big([(\boldsymbol{\theta}_n^{\mathcal{A}})^{\mathsf{T}},(\boldsymbol{\theta}_n^{\mathcal{A}^c})^{\mathsf{T}}]^{\mathsf{T}}\big) < \boldsymbol{\mathcal{L}}_n\big([(\boldsymbol{\theta}_n^{\mathcal{A}})^{\mathsf{T}},\boldsymbol{0}^{\mathsf{T}}]^{\mathsf{T}}\big),$$

where  $\boldsymbol{\theta}^{\star} = [(\boldsymbol{\theta}^{\star \mathcal{A}})^{\mathsf{T}}, (\boldsymbol{\theta}^{\star \mathcal{A}^{c}})^{\mathsf{T}}]^{\mathsf{T}}$ , with  $\mathcal{A}$  containing the indices of all nonzero elements of  $\boldsymbol{\theta}^{\star}$ .

Proof. Consider,

$$\begin{pmatrix} \mathcal{L}_n \left( [(\boldsymbol{\theta}_n^{\mathcal{A}})^{\mathsf{T}}, (\boldsymbol{\theta}_n^{\mathcal{A}^c})^{\mathsf{T}}]^{\mathsf{T}} \right) - \mathcal{L}_n \left( [(\boldsymbol{\theta}_n^{\mathcal{A}})^{\mathsf{T}}, \mathbf{0}^{\mathsf{T}}]^{\mathsf{T}} \right) \end{pmatrix} \\ = \frac{1}{(n-p)} \left( l_n(\boldsymbol{\theta}_n) - l_n \left( [(\boldsymbol{\theta}_n^{\mathcal{A}})^{\mathsf{T}}, \mathbf{0}^{\mathsf{T}}]^{\mathsf{T}} \right) \right) - \left( \sum_{k \in \mathcal{E}} r_{\lambda_n} \left( \theta_{n,k} \right) - \sum_{k \in \mathcal{E} \cap \mathcal{A}} r_{\lambda_n} \left( \theta_{n,k} \right) \right)$$

By the second-order Taylor expansion, we have

$$l_n(oldsymbol{ heta}_n) - l_n(oldsymbol{ heta}^{\star}) = ig(oldsymbol{ heta}_n - oldsymbol{ heta}^{\star}ig)^{\mathsf{T}} 
abla_{ heta} l_n(oldsymbol{ heta}_n) - l_n(oldsymbol{ heta}^{\star}) = ig([(oldsymbol{ heta}_n)^{\mathsf{T}}, oldsymbol{0}^{\mathsf{T}}]^{\mathsf{T}} - oldsymbol{ heta}^{\star}ig)^{\mathsf{T}} 
abla_{ heta} l_n(oldsymbol{ heta}_n)^{\mathsf{T}}, oldsymbol{0}^{\mathsf{T}}]^{\mathsf{T}} - oldsymbol{ heta}^{\star}ig)^{\mathsf{T}} 
abla_{ heta} l_n(oldsymbol{ heta}_n)^{\mathsf{T}}, oldsymbol{0}^{\mathsf{T}}]^{\mathsf{T}} - oldsymbol{ heta}^{\star}ig)^{\mathsf{T}} 
abla_{ heta} l_n(oldsymbol{ heta}_n)^{\mathsf{T}} 
abla_{ heta} l_n(oldsymbol{ heta}^{\star}) + rac{1}{2} ig([(oldsymbol{ heta}_n)^{\mathsf{T}}, oldsymbol{0}^{\mathsf{T}}]^{\mathsf{T}} - oldsymbol{ heta}^{\star}ig)^{\mathsf{T}} 
abla_{ heta} l_n(oldsymbol{ heta}_n)^{\mathsf{T}} 
abla_{ heta} l_n(oldsymbol{ heta}_n)^{\mathsf{T}} 
abla_{ heta} l_n(oldsymbol{ heta}^{\star}) + rac{1}{2} ig([(oldsymbol{ heta}_n)^{\mathsf{T}}, oldsymbol{0}^{\mathsf{T}}]^{\mathsf{T}} - oldsymbol{ heta}^{\star}ig)^{\mathsf{T}} 
abla_{ heta} l_n(oldsymbol{ heta}_n)^{\mathsf{T}} 
abla_{ heta} l_n(oldsymbol{ heta}_$$

In the above  $\tilde{\boldsymbol{\theta}}_n$  is  $s\boldsymbol{\theta}^{\star} + (1-s)\boldsymbol{\theta}_n$  for some 0 < s < 1 and  $\tilde{\boldsymbol{\theta}}_n^{\mathcal{A}}$  is  $t\boldsymbol{\theta}^{\star} + (1-t)[(\boldsymbol{\theta}_n^{\mathcal{A}})^{\mathsf{T}}, \mathbf{0}^{\mathsf{T}}]^{\mathsf{T}}$  for some 0 < t < 1. We know that  $\boldsymbol{\theta}^{\star} = [(\boldsymbol{\theta}^{\star \mathcal{A}})^{\mathsf{T}}, \mathbf{0}^{\mathsf{T}}]^{\mathsf{T}}$ , which gives

$$\left(\boldsymbol{\theta}_n - \boldsymbol{\theta}^\star\right)^\mathsf{T} 
abla_{\theta} l_n(\boldsymbol{\theta}^\star) = \left(\boldsymbol{\theta}_n^\mathsf{A} - \boldsymbol{\theta}^{\star\mathsf{A}}\right)^\mathsf{T} 
abla_{\theta^{\mathcal{A}}} l_n(\boldsymbol{\theta}^\star) + \left(\boldsymbol{\theta}_n^{\mathsf{A}^c} - \boldsymbol{\theta}^{\star\mathsf{A}^c}\right)^\mathsf{T} 
abla_{\theta^{\mathcal{A}^c}} l_n(\boldsymbol{\theta}^\star).$$

From Proposition 6.2(i) we have  $\nabla_{\theta} l_n(\theta^{\star}) = O_P(n^{1/2})$ . Then,

$$(\boldsymbol{\theta}_n - \boldsymbol{\theta}^{\star})^{\mathsf{T}} \nabla_{\boldsymbol{\theta}} l_n(\boldsymbol{\theta}^{\star}) = (\boldsymbol{\theta}_n^{\mathsf{A}} - \boldsymbol{\theta}^{\star \mathsf{A}})^{\mathsf{T}} \nabla_{\boldsymbol{\theta}^{\mathcal{A}}} l_n(\boldsymbol{\theta}^{\star}) + \mathrm{O}_{\mathrm{P}}(n^{1/2}) |\boldsymbol{\theta}_n^{\mathcal{A}^c}|$$

As  $n \to \infty$ ,  $\tilde{\boldsymbol{\theta}}_n \to \boldsymbol{\theta}^{\star}$  a.s., then from Proposition 6.2(ii) we have  $-n^{-1}\nabla_{\boldsymbol{\theta}}^2 l_n(\tilde{\boldsymbol{\theta}}_n) = \mathrm{I}(\boldsymbol{\theta}^{\star})(1+o_{\mathrm{P}}(1))$ . Furthermore, as  $n \to \infty$ ,  $\tilde{\boldsymbol{\theta}}_n^{\mathcal{A}} \to \boldsymbol{\theta}^{\star \mathcal{A}}$  a.s., we have  $-n^{-1}\nabla_{\boldsymbol{\theta}}^2 l_n(\tilde{\boldsymbol{\theta}}_n^{\mathcal{A}}) = \mathrm{I}_{11}(\boldsymbol{\theta}^{\star})(1+o_{\mathrm{P}}(1))$ , where the Fisher information matrix corresponding to  $\boldsymbol{\theta}^{\star} = [(\boldsymbol{\theta}^{\star \mathcal{A}})^{\mathsf{T}}, (\boldsymbol{\theta}^{\star \mathcal{A}^c})^{\mathsf{T}}]^{\mathsf{T}}$ , is written as

$$\mathrm{I}(oldsymbol{ heta}^{\star}) = egin{bmatrix} \mathrm{I}_{11}(oldsymbol{ heta}^{\star}) & \mathrm{I}_{12}(oldsymbol{ heta}^{\star}) \ \mathrm{I}_{12}(oldsymbol{ heta}^{\star}) & \mathrm{I}_{22}(oldsymbol{ heta}^{\star}) \end{bmatrix}.$$

Thus, for large enough n, we have

$$l_{n}(\boldsymbol{\theta}_{n}) - l_{n}\left([(\boldsymbol{\theta}_{n}^{\mathcal{A}})^{\mathsf{T}}, \boldsymbol{0}^{\mathsf{T}}]^{\mathsf{T}}\right) = O_{P}(n^{1/2})|\boldsymbol{\theta}_{n}^{\mathcal{A}^{c}}| - \frac{1}{2}n\left(\boldsymbol{\theta}_{n} - \boldsymbol{\theta}^{\star}\right)^{\mathsf{T}}I(\boldsymbol{\theta}^{\star})\left(\boldsymbol{\theta}_{n} - \boldsymbol{\theta}^{\star}\right)(1 + o_{P}(1))$$
$$+ \frac{1}{2}n\left(\boldsymbol{\theta}_{n}^{\mathcal{A}} - \boldsymbol{\theta}^{\star\mathcal{A}}\right)^{\mathsf{T}}I_{11}(\boldsymbol{\theta}^{\star})\left(\boldsymbol{\theta}_{n}^{\mathcal{A}} - \boldsymbol{\theta}^{\star\mathcal{A}}\right)(1 + o_{P}(1))$$
$$= O_{P}(n^{1/2})|\boldsymbol{\theta}_{n}^{\mathcal{A}^{c}}| - n\left(\left(\boldsymbol{\theta}_{n}^{\mathcal{A}} - \boldsymbol{\theta}^{\star\mathcal{A}}\right)^{\mathsf{T}}I_{12}(\boldsymbol{\theta}^{\star})\boldsymbol{\theta}_{n}^{\mathcal{A}^{c}}$$
$$+ \frac{1}{2}(\boldsymbol{\theta}_{n}^{\mathcal{A}^{c}})^{\mathsf{T}}I_{22}(\boldsymbol{\theta}^{\star})\boldsymbol{\theta}_{n}^{\mathcal{A}^{c}}\right)(1 + o_{P}(1)).$$

Since  $\|\boldsymbol{\theta}_n^{\mathcal{A}} - \boldsymbol{\theta}^{\star \mathcal{A}}\| = \mathcal{O}_{\mathcal{P}}(n^{-1/2})$  and  $\|\boldsymbol{\theta}_n^{\mathcal{A}^c}\| = \mathcal{O}_{\mathcal{P}}(n^{-1/2})$ , for large enough n and some constants  $\mathcal{C}_1, \mathcal{C}_2, \mathcal{C}_3 > 0$ , we get

$$l_{n}(\boldsymbol{\theta}_{n}) - l_{n}([(\boldsymbol{\theta}_{n}^{\mathcal{A}})^{\mathsf{T}}, \mathbf{0}^{\mathsf{T}}]^{\mathsf{T}}) \leq C_{1}n^{1/2}|\boldsymbol{\theta}_{n}^{\mathcal{A}^{c}}| + nC_{2}n^{-1/2}|\boldsymbol{\theta}_{n}^{\mathcal{A}^{c}}|(1 + o_{\mathrm{P}}(1)) \leq n^{1/2}C_{3}\sum_{k \in \mathcal{A}^{c}}|\boldsymbol{\theta}_{n,k}|.(16)$$

Next, we notice that

$$\left(\sum_{k\in\mathcal{E}}r_{\lambda_n}(\theta_{n,k})-\sum_{k\in\mathcal{E}\cap\mathcal{A}}r_{\lambda_n}(\theta_{n,k})\right)=\sum_{k\in\mathcal{E}\cap\mathcal{A}^c}r_{\lambda_n}(\theta_{n,k}).$$

From (16), for large enough n, we have

$$\mathcal{L}_n\big([(\boldsymbol{\theta}_n^{\mathcal{A}})^{\mathsf{T}}, (\boldsymbol{\theta}_n^{\mathcal{A}^c})^{\mathsf{T}}]^{\mathsf{T}}\big) - \mathcal{L}_n\big([(\boldsymbol{\theta}_n^{\mathcal{A}})^{\mathsf{T}}, \mathbf{0}^{\mathsf{T}}]^{\mathsf{T}}\big) \le \frac{n^{1/2}}{(n-p)} \mathcal{C}_3 \sum_{k \in \mathcal{A}^c} |\boldsymbol{\theta}_{n,k}| - \frac{1}{n} \sum_{k \in \mathcal{E} \cap \mathcal{A}^c} nr_{\lambda_n}(\boldsymbol{\theta}_{n,k}).$$
(17)

As  $\|\boldsymbol{\theta}_n - \boldsymbol{\theta}^{\star}\| = O_P(n^{-1/2})$ , for the coefficients belonging to the zero-elements set  $\mathcal{A}^c$ , we have  $|\boldsymbol{\theta}_{n,k}| = O_P(n^{-1/2})$ . Then, for each  $k \in \mathcal{A}^c$ , Assumption 3 is applicable on the penalty function value  $r_{\lambda_n}(\boldsymbol{\theta}_{n,k})$ , which gives

$$\liminf_{n\to\infty} nr_{\lambda_n}(\theta_{n,k}) = \liminf_{n\to\infty} \sqrt{n} |\theta_{n,k}| \sqrt{n} r'_{\lambda_n}(\theta_{n,k}) = +\infty \text{ a.s.}.$$

Hence, as  $n \to \infty$ , from (17) we deduce that

$$\mathcal{L}_nig([(oldsymbol{ heta}_n)^{\mathsf{T}},(oldsymbol{ heta}_n^{\mathcal{A}^c})^{\mathsf{T}}]^{\mathsf{T}}ig) - \mathcal{L}_nig([(oldsymbol{ heta}_n)^{\mathsf{T}},oldsymbol{0}^{\mathsf{T}}]^{\mathsf{T}}ig) < 0 \,\, \mathrm{a.s.}.$$

which proves the result.

Now, we provide the remainder of the proof. From Lemma 6.3, we know that for any  $\sqrt{n}$ -consistent estimator  $\widehat{\theta}_n$ , the parameter vector  $\begin{bmatrix} \widehat{\theta}_n^{\mathcal{A}} \\ 0 \end{bmatrix}$  maximizes  $\mathcal{L}_n(\theta)$  with probability tending to one as  $n \to \infty$ , over any other choice  $\theta_n = \begin{bmatrix} \theta_n^{\mathcal{A}} \\ \theta_n^{\mathcal{A}^c} \end{bmatrix}$  such that  $\|\theta_n - \theta^\star\| = O_P(n^{-1/2})$ . From Theorem 1, we know that a  $\sqrt{n}$ -consistent maximizer  $\widehat{\theta}_n$  of  $\mathcal{L}_n(\theta)$  exists when  $\lambda_n$  and  $r_{\lambda_n}$  satisfy Assumption 2. Then, it must be true that  $\mathbb{P}(\widehat{\theta}_n^{\mathcal{A}^c} = 0) \to 1$  as  $n \to \infty$ .

The result in Theorem 1 shows that we have a  $\sqrt{n}$ -consistent estimator if  $a_n = O(n^{-1/2})$ , with  $\lambda_n \to 0$ . For SCAD and MCP penalty functions, it is then enough that  $\lambda_n \to 0$  as  $n \to \infty$ , since then  $a_n = 0$ . For the LASSO penalty, we must have at least that  $\lambda_n = O(n^{-1/2})$  since  $a_n = \lambda_n$ ; and for

the ADALASSO, we require  $\sqrt{n\lambda_n} = o(1)$ . However, as is well-known, it is not possible to recover the sparsity structure using the LASSO penalty as we will then require that  $\lim_{n\to\infty} \sqrt{n\lambda_n} = \infty$ , to satisfy Assumption 3, which contradicts the earlier requirement on the tuning parameter decay rate for the LASSO penalty function. The sparsity structure can be recovered with the other three penalty functions, since we can choose  $\lambda_n \sim n^{-1/2-\psi}$  for a  $0 < \psi < \gamma/2$ , for the adaptive LASSO, and  $\lambda_n \sim n^{-1/2} \log n$  for the SCAD and MCP penalties.

If M is under-specified, the MPLE  $\hat{\theta}_n$  converges to the minimizer of the Kullback-Leibler distance between the densities of the under-specified and true models, a property shared by the MLE under this misspecification (Douc and Moulines, 2012). On the other hand, in Theorem 2 we show that if M is over-specified, the density function of the over-fitted MSVAR model based on MPLE consistently estimates the density function of the true model, which is useful for prediction. In particular, we show that the estimated predictive density of the over-fitted model based on MPLE consistently estimates the *h*-step ahead predictive density of the true MSVAR model with M regimes, denoted by  $f^*(\mathbf{y}_{n+1:h}|\mathbf{y}_{1:n})$ , and computed in Section 4 of the main manuscript. As a consequence of this result, in practice when the true number of regimes is unknown, a conservative choice of M considering the sample size *n* can guarantee a reliable estimate of the *h*-step ahead predictive density, and the optimal predictor (see Section 4, main manuscript) in the sense of minimum mean squared prediction error.

**Theorem 2.** Under Assumption 1–3, the estimated h-step ahead predictive density function  $\widehat{f}_{\mathcal{M}}(\mathbf{y}_{n+1:h}|\mathbf{y}_{1:n})$  for number of regimes  $\mathcal{M} \geq \mathcal{M}$ , converges almost surely to the true h-step ahead predictive density  $f^{\star}(\mathbf{y}_{n+1:h}|\mathbf{y}_{1:n})$  as  $n \to \infty$  where  $\mathcal{M}$  is the true number of regimes in the model.

*Proof.* We provide the proof for the one-step estimated predictive density. The result for the *h*-step estimated predictive density can be shown similarly.

We denote the generic intercept vector for VAR model by  $\nu$  and the vectorized VAR coefficients and covariance matrices as  $a_i = \mathbf{vec}(A_i), i = 1, ..., p$  and  $\sigma = \mathbf{vec}(\Sigma)$ . For number of regimes  $\mathcal{M} > \mathcal{M}$  and n > p, the one-step ahead predictive density function is given as

$$f_{\mathcal{M}}(\mathbf{y}_{n+1}|\mathbf{y}_{1:n}) = \sum_{m=1}^{\mathcal{M}} g(\mathbf{y}_{n+1}, \mathbf{y}_{1:n}; \nu^{(m)}, \{a_i^{(m)}\}_{i=1}^p, \sigma^{(m)}) \mathbb{P}(s_{n+1} = m|\mathbf{y}_{1:n}),$$

where

$$g(\mathbf{y}_{n}+1,\mathbf{y}_{1:n};\nu,\{a_{i}\}_{i=1}^{p},\sigma) := \frac{1}{\sqrt{(2\pi)^{d}|\Sigma|}} \exp\left(-\frac{1}{2}(\mathbf{y}_{n+1}-\mu(\mathbf{y}_{1:n},\nu,\{a_{i}\}_{i=1}^{p}))^{\mathsf{T}}(\Sigma)^{-1}\right)$$
$$(\mathbf{y}_{n+1}-\mu(\mathbf{y}_{1:n},\nu,\{a_{i}\}_{i=1}^{p})),$$
$$\mu(\mathbf{y}_{1:n},\nu,\{a_{i}\}_{i=1}^{p}) := \nu + A_{1}\mathbf{y}_{n} + \ldots + A_{p}\mathbf{y}_{n-p+1}.$$

The estimated one-step ahead predictive density for number of regimes  $\mathcal{M}$  is given as

$$\widehat{f}_{\mathcal{M}}(\mathbf{y}_{n+1}|\mathbf{y}_{1:n}) = \sum_{j=1}^{\mathcal{M}} g\big(\mathbf{y}_{n+1}, \mathbf{y}_{1:n}; \widehat{\nu}^{(m)}, \{\widehat{a}_{i}^{(m)}\}_{i=1}^{p}, \widehat{\sigma}^{(m)}\big) \widehat{\mathbb{P}}(s_{n+1} = m | \mathbf{y}_{1:n}),$$

with  $\mathbb{P}(s_{n+1} = m | \mathbf{y}_{1:n})$  computed via forward and backward recursions and using the estimates of the transition probability matrix for  $\mathcal{M}$  number of regimes. The estimates of VAR coefficients and covariance matrices are denoted by  $(\hat{\nu}^{(m)}, \{\hat{a}_i^{(m)}\}_{i=1}^p)$  and  $\hat{\sigma}^{(m)}$  respectively. We can alternatively write

$$\widehat{f}_{\mathcal{M}}(\mathbf{y}_{n+1}|\mathbf{y}_{1:n}) = \int g(\mathbf{y}_{n+1}, \mathbf{y}_{1:n}; \nu, \{a_i\}_{i=1}^p, \sigma) \mathrm{d}\widehat{\mathbf{\Phi}}_{\mathcal{M}, n}(\nu, \{a_i\}_{i=1}^p, \sigma),$$
(18)

with

$$\widehat{\boldsymbol{\Phi}}_{\mathcal{M},n}(\nu, \{a_i\}_{i=1}^p, \sigma) = \sum_{m=1}^{\mathcal{M}} \widehat{\mathbb{P}}(s_{n+1} = m | \mathbf{y}_{1:n}) \mathrm{H}(\nu \ge \widehat{\nu}^{(m)}, \{a_i\}_{i=1}^p \ge \{\widehat{a}_i^{(m)}\}_{i=1}^p, \sigma \ge \widehat{\sigma}^{(m)}),$$

where  $H(\cdot)$  denotes the Heaviside step function. The true one-step ahead predictive density can be then written as

$$f^{\star}(\mathbf{y}_{n+1}|\mathbf{y}_{1:n}) = \int g(\mathbf{y}_{n+1}, \mathbf{y}_{1:n}; \nu, \{a_i\}_{i=1}^p, \sigma) \mathrm{d}\mathbf{\Phi}_n^{\star}(\nu, \{a_i\}_{i=1}^p, \sigma),$$

with

$$\Phi_n^{\star}(\nu, \{a_i\}_{i=1}^p, \sigma) = \sum_{m=1}^{M} \mathbb{P}^{\star}(s_{n+1} = m | \mathbf{y}_{1:n}) \mathbf{H}(\nu \ge \nu^{(m)^{\star}}, \{a_i\}_{i=1}^p \ge \{a^{(m)^{\star}}_i\}_{i=1}^p, \sigma \ge \sigma^{(m)^{\star}}).$$

The estimated and true filtered probabilities are computed by fixing the initial state  $s_p$  and initializing with true conditional state probabilities  $\mathbb{P}^{\star}(s_p = m | \mathbf{y}_{1:p})$  respectively. The effect of initial distribution dissipates geometrically fast in number of observations n as shown in Douc et al. (2004, Corollary 1). Next, we define the following distance between the probability measures  $\Phi_n^{\star}(\nu, \{a_i\}_{i=1}^p, \sigma)$ and  $\Phi_{\mathcal{M},n}(\nu, \{a_i\}_{i=1}^p, \sigma)$ :

$$D(\boldsymbol{\Phi}_{\mathcal{M},n}, \boldsymbol{\Phi}_{n}^{\star})$$

$$= \int \int \left| \boldsymbol{\Phi}_{\mathcal{M},n}(\nu, \{a_{i}\}_{i=1}^{p}, \sigma) - \boldsymbol{\Phi}_{n}^{\star}(\nu, \{a_{i}\}_{i=1}^{p}, \sigma) \right| e^{-(|\nu| + \sum_{i=1}^{p} |a_{i}| + |\sigma|)} d\nu da_{1} \dots da_{p} d\sigma d\mathbb{P}^{\star}(\mathbf{y}_{1:n}).$$

$$(19)$$

The above distance metrizes the space of probability measures on the space of VAR model parameter  $(\nu, \{a_i\}_{i=1}^p, \sigma) \subset \Theta$ . We can consider any other distance between probability measures on the space of VAR model parameter  $(\nu, \{a_i\}_{i=1}^p, \sigma)$ . For an arbitrary  $\delta > 0$ , we consider a family

$$\mathfrak{H}_n(\delta) = \{ \mathbf{\Phi}_{\mathfrak{M},n} : \mathrm{D}(\mathbf{\Phi}_{\mathfrak{M},n},\mathbf{\Phi}_n^{\star}) > \delta \},$$

of probability measures such that each element in it is at least a  $\delta$  distance away from  $\Phi_n^{\star}$ . Then, clearly  $\Phi_n^{\star} \notin \mathcal{H}_n(\delta)$ , and

$$\mathbb{E}^{\star}\left[\log\left(\frac{f_{\mathcal{M}}(\mathbf{y}_{n+1}|\mathbf{y}_{1:n})}{f^{\star}(\mathbf{y}_{n+1}|\mathbf{y}_{1:n})}\right)\right] < 0$$

Similarly, we also have for  $t = p + 1, \ldots, n$ 

$$\mathbb{E}^{\star}\left[\log\left(\frac{f_{\mathcal{M}}(\mathbf{y}_t|=\mathbf{y}_{1:t-1})}{f^{\star}(\mathbf{y}_t|\mathbf{y}_{1:t-1})}\right)\right] < 0,$$

where the conditional densities  $f_{\mathcal{M}}(\mathbf{y}_t|\mathbf{y}_{1:t-1})$  and  $f^{\star}(\mathbf{y}_t|\mathbf{y}_{1:t-1})$  have their own representation similar to  $f_{\mathcal{M}}(\mathbf{y}_{n+1}|\mathbf{y}_{1:n})$  and  $f^{\star}(\mathbf{y}_{n+1}|\mathbf{y}_{1:n})$  but with their own probability measures  $\Phi_{\mathcal{M},t} \in \mathcal{H}_t(\delta)$  and  $\Phi_t^{\star}$ , respectively. From the stationarity of the process and ergodic theorem, we can also conclude that

$$\frac{1}{n-p}\sum_{t=p+1}^{n}\log\left(\frac{f_{\mathcal{M}}(\mathbf{y}_{t}|\mathbf{y}_{1:t-1})}{f^{\star}(\mathbf{y}_{t}|\mathbf{y}_{1:t-1})}\right) = \frac{1}{n-p}\log\left(\frac{f_{\mathcal{M}}(\mathbf{y}_{p+1},\dots,\mathbf{y}_{n}|\mathbf{y}_{1:p})}{f^{\star}(\mathbf{y}_{p+1},\dots,\mathbf{y}_{n}|\mathbf{y}_{1:p})}\right) < -\epsilon(\delta) \text{ a.s.}, \quad (20)$$

for some  $\epsilon(\delta) > 0$  and *n* large enough. Define  $\gamma := (\nu, \{a_i\}_{i=1}^p, \sigma)$  as the VAR model parameter vector. Then, the joint probability density under the assumption of  $\mathcal{M} > \mathcal{M}$  regimes and true joint probability density can be expressed as follows:

$$f_{\mathcal{M}}(\mathbf{y}_{p+1},\ldots,\mathbf{y}_{n}|\mathbf{y}_{1:p}) = \int \prod_{t=p+1}^{n} g(\mathbf{y}_{t},\mathbf{y}_{1:t-1};\boldsymbol{\gamma}_{t}) \mathrm{d}\mathbf{\Phi}_{\mathrm{joint},\mathcal{M}}(\boldsymbol{\gamma}_{p+1:n})$$
$$f^{\star}(\mathbf{y}_{p+1},\ldots,\mathbf{y}_{n}|\mathbf{y}_{1:p}) = \int \prod_{t=p+1}^{n} g(\mathbf{y}_{t},\mathbf{y}_{1:t-1};\boldsymbol{\gamma}_{t}) \mathrm{d}\mathbf{\Phi}^{\star}_{\mathrm{joint}}(\boldsymbol{\gamma}_{p+1:n}),$$

where  $\gamma_{p+1:n} = (\gamma_{p+1}, \ldots, \gamma_n)$ , with  $\gamma_t, t = p+1, \ldots, n$ , representing the choice of VAR model parameter at different instances. Furthermore,

$$g(\mathbf{y}_{t}, \mathbf{y}_{1:t-1}; \boldsymbol{\gamma}) := \frac{1}{\sqrt{(2\pi)^{d} |\Sigma|}} \exp\left(-\frac{1}{2}(\mathbf{y}_{t} - \mu(\mathbf{y}_{1:t-1}, \nu, \{a_{i}\}_{i=1}^{p}))^{\mathsf{T}}(\Sigma)^{-1}(\mathbf{y}_{t} - \mu(\mathbf{y}_{1:t-1}, \nu, \{a_{i}\}_{i=1}^{p}))\right),$$
$$\mu(\mathbf{y}_{1:t-1}, \nu, \{a_{i}\}_{i=1}^{p}) = \nu + A_{1}\mathbf{y}_{t-1} + \ldots + A_{p}\mathbf{y}_{t-p},$$

and

$$\Phi_{\text{joint},\mathcal{M}}(\boldsymbol{\gamma}_{p+1:n}) = \sum_{j_{p+1}=1}^{\mathcal{M}} \dots \sum_{j_{n}=1}^{\mathcal{M}} \mathbb{P}(s_{p+1} = j_{p+1} | \mathbf{y}_{1:p}) \prod_{t=p+2}^{n} \alpha'_{j_{t-1},j_{t}} \prod_{t=p+1}^{n} \mathrm{H}(\gamma_{t} \ge \gamma'_{j_{t}}),$$

$$\Phi_{\text{joint}}^{\star}(\boldsymbol{\gamma}_{p+1:n}) = \sum_{j_{p+1}=1}^{\mathcal{M}} \dots \sum_{j_{n}=1}^{\mathcal{M}} \mathbb{P}^{\star}(s_{p+1} = j_{p+1} | \mathbf{y}_{1:p}) \prod_{t=p+2}^{n} \alpha_{j_{t-1},j_{t}}^{\star} \prod_{t=p+1}^{n} \mathrm{H}(\gamma_{t} \ge \gamma'_{j_{t}}).$$

For the joint distribution  $\Phi_{\text{joint},\mathcal{M}}(\gamma_{p+1:n})$  we can similarly define a family

$$\mathfrak{H}_{ ext{joint}}(\delta) := \{ \mathbf{\Phi}_{ ext{joint}, \mathcal{M}} : \mathrm{D}(\mathbf{\Phi}_{ ext{joint}, \mathcal{M}}, \mathbf{\Phi}^{\star}_{ ext{joint}}) > \delta \}$$

based on its distance from the joint distribution  $\Phi_{\text{joint}}^{\star}(\gamma_{p+1:n})$  with the distance being defined analogously as in (19) with inner integral being with respect to  $\gamma_{p+1:n}$  and the outside probability measure corresponding to  $\mathbb{P}^{\star}(\mathbf{y}_{1:p})$ .

From the result in (20) for any  $\Phi_{\text{joint},\mathcal{M}} \in \mathcal{H}_{\text{joint}}(\delta)$ , and the fact that penalty terms are non-negative with order o(1), we get

$$\sup_{\mathcal{H}_{\text{joint}}(\delta)} \left( \frac{1}{(n-p)} \log \left( \frac{f_{\mathcal{M}}(\mathbf{y}_{p+1}, \dots, \mathbf{y}_n | \mathbf{y}_{1:p})}{f^{\star}(\mathbf{y}_{p+1}, \dots, \mathbf{y}_n | \mathbf{y}_{1:p})} \right) - \left( \sum_{k \in \mathcal{E}_{\mathcal{M}}} \mathrm{R}_{\lambda_n} \left( \theta_k^{\mathcal{T}} \right) - \sum_{k \in \mathcal{E}} \mathrm{R}_{\lambda_n} \left( \theta_k^{\star} \right) \right) \right) < -\epsilon(\delta) \text{ a.s.}.$$

Hence, the joint probability distribution corresponding to the penalized maximum-likelihood estimate in coefficient space  $\Theta_{\mathcal{M}}$  cannot be an element of  $\mathcal{H}_{\text{joint}}(\delta)$  almost surely as  $n \to \infty$ . This is true for any  $\delta > 0$ , thus we must have for the joint distribution corresponding to the penalized maximum-likelihood estimate in coefficient space  $\Theta_{\mathcal{M}} : \widehat{\Phi}_{\text{joint},\mathcal{M}}$  that  $D(\widehat{\Phi}_{\text{joint},\mathcal{M}}, \Phi_{\text{joint}}^*) \to 0$  as  $n \to \infty$ . In other words, we have that  $\widehat{\Phi}_{\text{joint},\mathcal{M}}$  converges weakly to  $\Phi_{\text{joint}}^*$  as  $n \to \infty$ . Then we must also have that  $\widehat{\Phi}_{\mathcal{M},n}$  converges weakly to the true distribution  $\Phi_n^*$  as  $n \to \infty$ . As the function g in (18) is bounded and continuous for the choice of parameter  $\theta$  in the compact parameter space  $\Theta$ , we have that  $\widehat{f}_{\mathcal{M}}(\mathbf{y}_{n+1}|\mathbf{y}_{1:n})$  converges almost surely to  $f^*(\mathbf{y}_{n+1}|\mathbf{y}_{1:n})$  as  $n \to \infty$ .  $\Box$ 

# A3 Complementary numerical results

Recall the scenarios **S1** and **S2** for sparse covariance and precision matrices, respectively, along with sparse AR matrices. We use MLE\* to refer to the MLE obtained by incorporating the knowledge of the true zero parameters of an MSVAR model, which we denote it by  $\tilde{\theta}$  in the main paper. We set M = 2, p = 1, and consider dimensions d = 20 with sample sizes  $n \in \{200, 300, 400\}, d = 40$ with  $n \in \{300, 400, 500\}$ , and d = 100 with  $n \in \{600, 700, 800\}$ . For these values of (d, M, p), the parameter vector  $\theta$  has dimensions K = M $(d + pd^2 + d(d + 1)/2) + M(M-1) = 1262, 4922$  and 30302, respectively. The corresponding dimensions of the parameter vector of the true data-generating MSVAR models are 104, 202 and 504, respectively. The simulation result for each d should be analyzed on its own, since the parameter configurations of the underlying models corresponding to d = 20, 40, 100, are different; see Section 5 of the main manuscript on "Simulation design".

Sparsity scenario S1. Figure 1 shows the relative estimation error (REE) and true positive rate (TPR) for d = 20. The results for dimensions d = 40,100, are presented in the main manuscript. In Figure 1, we observe that in terms of the overall REE for the smallest sample size, SCAD and MCP perform better compared to the other two penalty functions. On the other hand, ADALASSO and MCP outperform when n = 300,400, while also being comparable to the MLE\* (median REE at most 1.2). In terms of TPR, the performance of the method based on all four penalties is reasonable.

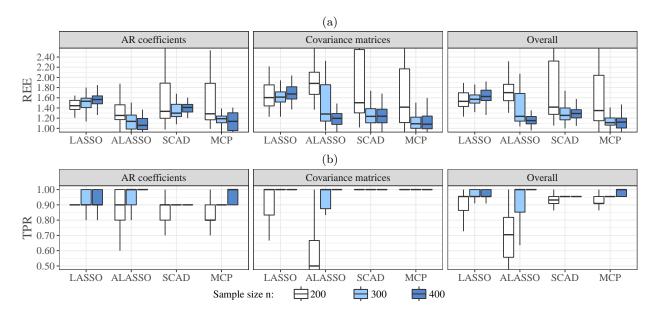


Figure 1: (a) Relative estimation error (REE) and (b) true positive rate (TPR, nonzero parameter detection, bottom) based on 50 random samples for data dimension d = 20, parameter dimension K = 1262, and sparsity scenario **S1**.

For d = 20, we also investigate the performance of the MPLE for a wider range of sample sizes  $n \in \{120, 200, 400, 800, 2000, 5000, 10000\}$ . Figure 2 shows the estimation error (EE) and the TPR values. For the sample sizes n = 120, 200, the results show that the EE are relatively large and the TPR are relatively low. This is expected as these sample sizes are close to the number of parameters in the true data-generating MSVAR model, which is 104. On the other hand, the results show that as the sample size increases to 400 and beyond the standard deviation of the estimates decreases, which is expected as per our result in Theorem 1-(i) on the consistency of the MPLE. In addition, the TPR reaches the value 1.0 as n increases, which confirms the sparsity recovery property of the

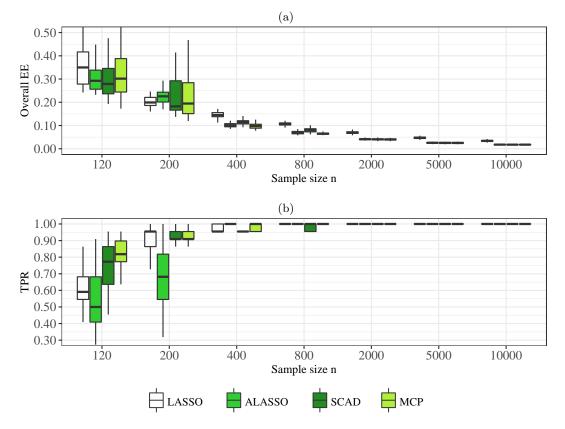


Figure 2: (a) Overall estimation error (EE) and (b) true positive rate (TPR) for data dimension d = 20, parameter dimension K = 1262, and sparsity scenario **S1**.

MPLE in Theorem 1-(ii). We also performed simulations for M = 2, p = 1 and d = 40 (K = 4922) with a minimum sample size n = 220 (the number of parameters in the true data-generating MSVAR model is 202), and we observed the same behaviour of the MPLE as for d = 20.

Sparsity scenario S2. Relative estimation errors (REE) with respect to the MLE\* for this scenario are presented in Figure 3 for d = 20, 40, 100. Different from scenario S1, here we observe that the MPLE attains a lower estimation error compared to the MLE\*  $\tilde{\theta}$ , the reason being as follows. To obtain  $\tilde{\theta}$  for the case of sparse precision matrices  $\Omega^{(m)}$ , we first compute the regime-specific MLE of  $\Sigma^{(m)}$  and we then set the entries of the MLE of  $(\Sigma^{(m)})^{-1}$  to zero, corresponding to the zero entries of the true precision matrices. This estimation procedure does not directly use the knowledge of the true zero parameters in the precision matrices resulting in REE < 1, that is, a higher estimation error (EE) for  $\tilde{\theta}$  compared to the proposed penalized estimators. From the results we can see that regarding the AR matrices, the ADALASSO outperforms the other penalties, followed by SCAD, MCP and LASSO. About the precision matrices and also the overall REE, the four penalties perform similar to each other.

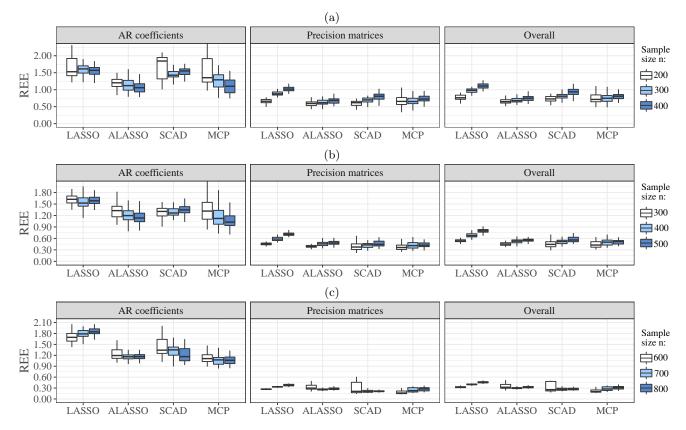
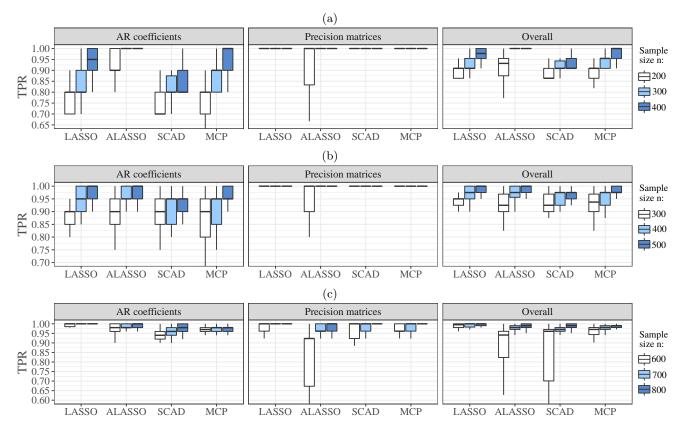


Figure 3: Relative estimation error (REE) based on 50 random samples for sparsity scenario **S2**: (a) d = 20, K = 1262 (b) d = 40, K = 4922 (c) d = 100, K = 30302, where d and K are the data and parameter dimensions, respectively.

Figure 4 shows the boxplots of the TPR. We can see that the method has a reasonable performance, above 0.85 overall. As in scenario **S1**, we observe mean true negative rate (TNR) above 0.90 for all cases (omitted).

Similar to scenario **S1**, for d = 20, we also investigate the performance of the MPLE for a wider range of sample sizes  $n \in \{120, 200, 400, 800, 2000, 5000, 10000\}$ , and for d = 40 with  $n \in \{220, 300, 400, 800, 2000, 5000, 10000\}$ . The performance of the method in terms of the overall EE



and TPR improves as the sample size increases (the results omitted), which is expected as per the result of Theorem 1.

Figure 4: True positive rate (TPR, nonzero parameter detection) based on 50 random samples for scenario **S2**: (a) d = 20, K = 1262 (b) d = 40, K = 4922 (c) d = 100, K = 30302, where d and K are the data and parameter dimensions, respectively.

**Diagonal covariance matrices.** In our simulation experiments considered in the main manuscript, we considered non-structured sparse covariance matrices. Here, we present the results for an experiment for d = 20, M = 2, p = 1 and diagonal true covariance matrices  $\Sigma^{(m)}$ . If we incorporate the knowledge of the true covariance matrices being diagonal, the number of parameters (dimension of  $\theta$ ) to be estimated by the MPLE reduces from K = 1262 to K = 882.

In Figure 5, corresponding to K = 1262, we observe that the median REE across all the four penalties is less than 1.6, with the ADALASSO and LASSO performing closer to the MLE\* compared to the other two penalties, as also shown by the overall EE. In terms of TPR and TNR, we observe that the MPLE is able to recover both the the true nonzero and zero (off-diagonal) entries of the covariance matrices, with median rates above 0.80 and 0.90, respectively.

In Figure 6, corresponding to K = 882, we observe that, by incorporating the knowledge of the covariance matrices being diagonal, the medians of all the performance measures remain roughly the same as in Figure 5. The main differences are that (i) the standard deviations of the REE and EE are lower in Figure 6 for the smallest sample size, and (ii) the TNR corresponding to the covariance matrices is now 1.0. This shows that the performance of our method is at par with the performance of the estimator that uses the knowledge of the true sparsity structure. In simulation designs with covariance matrices being multiples of the identity matrix, as well as for d = 40, we observe the same phenomenon (results omitted).

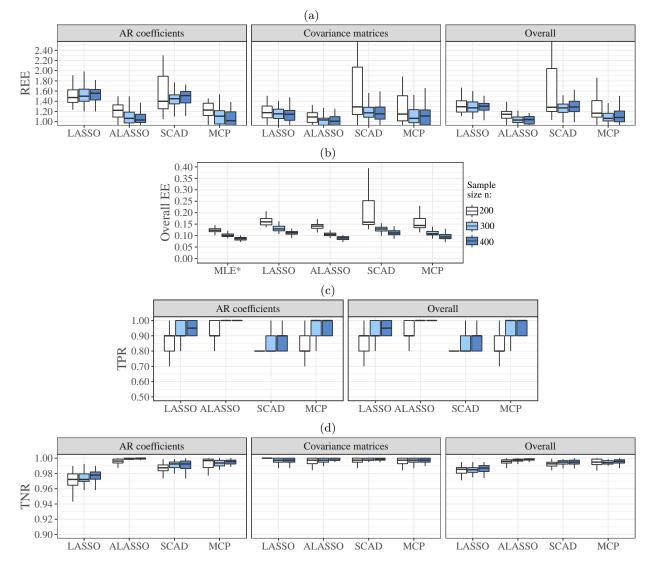


Figure 5: (a) Relative estimation error (REE), (b) overall estimation error (EE), (c) true positive rate (TPR) and (d) true negative rate (TNR) for model with diagonal covariance matrices and data dimension d = 20 (parameter dimension K = 1262). MLE\* represents maximum likelihood estimation knowing the location of the zero parameters.

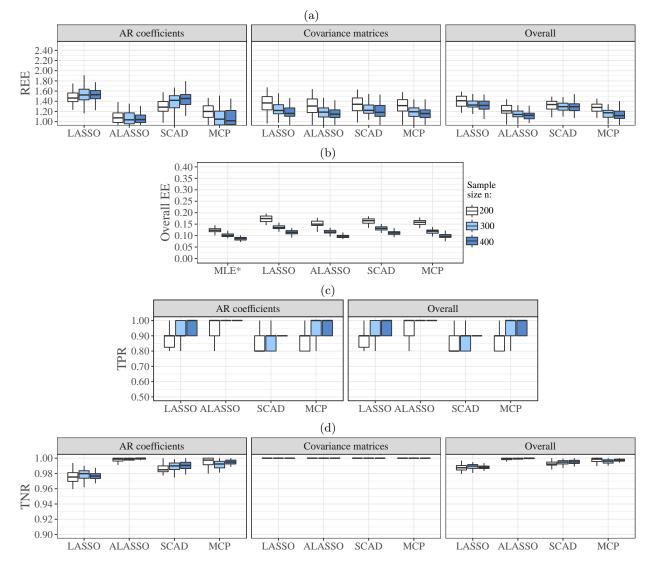


Figure 6: (a) Relative estimation error (REE), (b) overall estimation error (EE), (c) true positive rate (TPR) and (d) true negative rate (TNR) for model with diagonal covariance matrices and data dimension d = 20, using the knowledge of the true covariance matrices being diagonal (parameter dimension K = 882). MLE\* represents maximum likelihood estimation knowing the location of the zero parameters.

### A4 Model fitting results for the macroeconomic dataset case study

In the case study based on the Canadian macroeconomic dataset as presented in Section 6 of the main manuscript, we fit MSVAR models with M = 1, 2, 3, 4, and p = 1, 2, 3, and for each penalty and sparsity scenario **S1** (sparse covariance matrices) and scenario **S2** (sparse precision matrices). We compute BIC and MDL to choose the final model. From Table 1 below, we conclude that: (i) scenario **S1** yields the lowest values of BIC in all cases, and of MDL in almost all cases, compared to **S2**; (ii) under **S1** and for any penalty, the number of regimes M = 3 is consistently selected based on BIC and MDL; (iii) under **S1** and for M = 3, the ADALASSO and MCP provide the lowest values of both BIC and MDL. The change in BIC and MDL when varying M from 3 to 4 is not significant. Moreover, the optimization problems associated to  $M \ge 4$  are obviously more sensitive to their initialization, and more prone to bad local optima.

We employ our proposed AR-order selection method based on the hierarchical group-LASSO penalized estimation (Section A1.6), and obtain  $\hat{p} = 1$  for the estimated  $\hat{M} = 3$ . Finally, regarding the choice of the penalty function, since we are also interested in the predictive performance of the model, we choose between ADALASSO and MCP by comparing the value of their predictive densities obtained with  $\hat{M} = 3$  and  $\hat{p} = 1$ , evaluated on out-of-sample observations from the last 6 months in the dataset. The ADALASSO penalty yields the highest value of the log-predictive density (-162.4) compared to LASSO (-180.2), SCAD (-218.4) and MCP (-220.5). Thus, using ADALASSO , we obtain the final model with  $\hat{M} = 3$ ,  $\hat{p} = 1$ .

LASSO		M = 1		M = 2		M = 3		$\mathrm{M}=4$	
		BIC	MDL	BIC	MDL	BIC	MDL	BIC	MDL
<b>S</b> 1	p = 1	12224.3	-	11319.8	11219.4	11273.4	11096.9	11356.2	11170.5
	p=2	12095.8	-	11246.8	11158.9	11103.9	10959.7	11174.1	10960.8
	p=3	12049.2	-	11341.5	11257.3	11182.6	10650.2	11218.1	11034.4
<b>S2</b>	p = 1	12285.7	-	11938.6	11824.6	12123.4	11744.9	12071.9	11796.1
	p=2	12161.2	-	11822.4	11697.3	12507.9	11968.1	12149.1	11766.7
	p=3	12119.9	-	12061.6	11992.6	11778.0	11632.3	11958.6	11691.6
ALASSO		M =	1	M = 2		M = 3		M = 4	
		BIC	MDL	BIC	MDL	BIC	MDL		MDL
<b>S</b> 1	p = 1	12094.5	_	11051.7	10974.2	11009.9	10861.7	11095.1	10769.3
	p = 2	11931.5	_	10912.0	10833.0	10893.1	10758.6	10957.2	10778.9
	p=3	11872.9	-	10937.1	10848.1	10976.1	10828.9	10945.6	10739.6
<b>S2</b>	p = 1	12246.8	-	12617.7	12024.7	12125.2	11577.5	12090.3	11580.3
	p=2	12111.0	-	11970.6	11826.1	12311.5	11941.4	12618.7	11765.5
	p=3	12045.9	_	11622.8	11507.8	11551.3	11377.3	11787.7	11484.0
SCAD		M = 1		M = 2		M = 3		M = 4	
00110		BIC	MDL	BIC	MDL	BIC	MDL	BIC	MDL
<b>S</b> 1	p = 1	12187.4	_	11137.5	11059.7	11105.5	10917.0	11115.4	10854.5
	p = 2	12053.8	_	11043.6	10968.5	10943.6	10788.9	11014.4	10783.0
	p=3	12019.9	_	11186.0	11070.8	11051.9	10914.7	11110.8	10896.7
S2	p = 1	12322.3	_	12676.2	12305.7	11761.1	11539.9	12598.5	11710.1
	p=2	12182.4	-	11948.9	11817.0	12169.0	11731.6	11997.9	11845.1
	p=3	12118.2	_	12671.9	12306.7	12394.5	10049.0	12862.4	12069.9
MCP	1	M = 1		M = 2		M = 3		M = 4	
		BIC	MDL	BIC	MDL	BIC	MDL	BIC	MDL
<b>S</b> 1	p = 1	12074.3	-	11016.8	10953.2	10987.6	10754.9	11387.9	10976.8
	p = 2	11897.1	-	10885.6	10809.5	10840.4	10644.4	10917.7	10551.2
	p = 3	11863.4	-	10905.2	10841.1	10757.4	10600.7	10982.2	10630.4
S2	p = 1	12239.9	-	12166.7	11903.8	11864.8	11486.9	12409.5	11759.5
	p=2	12063.3	-	11927.1	11691.2	12205.6	11746.1	12603.9	12006.7

Table 1: BIC and MDL values based on the macroeconomic dataset of Section 6 of the main manuscript, for M = 1, 2, 3, 4, p = 1, 2, 3, sparsity scenarios **S1** and **S2**, and all the penalties considered. Minima for fixed p and criterion are highlighted in boldface.

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