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Multivariate cluster-weighted models based on seemingly unrelated linear regression

Cecilia Diani, Giuliano Galimberti, Gabriele Soffritti*

Abstract

A class of cluster-weighted models for a vector of continuous random variables is proposed. This class provides an extension to cluster-weighted modelling of multivariate and correlated responses that let the researcher free to use a different vector of covariates for each response. The class also includes parsimonious models obtained by imposing suitable constraints on the component-covariance matrices of either the responses or the covariates. Conditions for model identifiability are illustrated and discussed. Maximum likelihood estimation is carried out by means of an expectation-conditional maximisation algorithm. The effective

tiveness and usefulness of the proposed models are shown through the analysis

of simulated and real datasets.

Keywords: cluster analysis, ECM algorithm, Gaussian mixture model,

multivariate linear regression, parsimonious model

2010 MSC: 62J05, 62H12, 62F12

1. Introduction

Cluster-weighted modelling is a flexible framework for data analysis intro-

duced by Gershenfeld (1997) in which the joint distribution of a given random

 $_4$ vector is modelled by assuming that this vector is composed of an outcome ${f Y}$

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(response, dependent variable) and its explanatory variables **X** (covariates, predictors); in order to account for the possible presence of unknown clusters of observations, a finite mixture is embedded into the model. Thus, cluster-weighted models are useful to perform multivariate regression analysis with random covariates in the presence of unobserved heterogeneity. Such models play a prominent role when the sample observations come from several sub-populations, the distribution of the outcome as well as the effect of the covariates on the response change with the sub-populations and the covariates are not under the control of the researcher.

An intense research into cluster-weighted models has been carried out over the last decade. Ingrassia et al. (2012) and Ingrassia et al. (2014) have developed 15 models for continuous variables under both Gaussian and Student t mixture distributions. Solutions suitable for dealing with various types of responses are detailed in Punzo and Ingrassia (2013), Punzo and Ingrassia (2015), Ingrassia et al. (2015) and Di Mari et al. (2020). Models with non-linear relationships or many covariates have been proposed by Punzo (2014) and Subedi et al. (2013), 20 respectively. Robustified solutions have been developed by Subedi et al. (2015) 21 and Punzo and McNicholas (2017). As far as vectors of continuous random variables with a multivariate response are concerned, Dang et al. (2017) have 23 developed a family of parsimonious Gaussian cluster-weighted models, where suitable constraints are imposed on the eigen-decomposition of the component-25 covariance matrices so as to mitigate the problem of a large number of model parameters when dealing with several variables. An underlying assumption in the family of parsimonious Gaussian cluster-weighted models introduced by Dang et al. (2017) is that all the covariates in the model affect each examined response. However, in some situations there may be prior information concerning 30 the absence of certain covariates from the linear term employed in the prediction of a certain response, and different covariates may be expected to be relevant in the prediction of different responses, as in the seemingly unrelated regression 33 context (Srivastava and Giles, 1987). This approach to multivariate regression has been extensively employed in the modelling of multivariate economic

data, where some given aspects of economic behaviour are typically assumed to depend on different economic variables according to a certain general theory. Classical examples can be found in White and Hewings (1982) and Giles and Hampton (1984), where multivariate regression models with different vectors of covariates were specified and estimated based on employment equations and Cobb-Douglas production functions in different geographical locations, respec-41 tively. Other fields in which the same approach has been successfully employed are medicine, food quality, tourism economics, quality of life and health (see, e.g., Keshavarzi et al., 2012; Cadavez and Henningsen, 2012; Keshavarzi et al., 2013; Disegna and Osti, 2016; Heidari et al., 2017). Other regression models for multivariate responses based on finite mixture models have been introduced by Soffritti and Galimberti (2011); Dang and McNicholas (2015); Galimberti et al. (2016). The flexmix package (Grün and Leisch, 2008) in the R environment (R Core Team, 2020) provides a general framework for the specification and estimation of finite mixtures of regression models. 50

This paper introduces a class of multivariate seemingly unrelated Gaussian 51 linear cluster-weighted models. Models from this class are able to capture both 52 the linear dependencies among responses and the linear effects of the covariates on the responses from sample observations coming from heterogeneous popu-54 lations. Furthermore, with these models the researcher is enabled to specify a different vector of covariates for each response. The paper addresses the 56 model identification and maximum likelihood (ML) estimation. This latter task is carried out by resorting to an expectation-conditional maximisation (ECM) algorithm. In order to keep the total number of parameters as low as possible, parsimonious models are included into the novel class, where parsimony is 60 attained by constraining the component-covariance matrices using a parameteri-61 sation for such matrices which is based on their spectral decomposition (see, e.g., Celeux and Govaert, 1995). With this approach, fourteen different covariance structures are allowed for both the covariates and the responses. The usefulness and the great flexibility of the resulting model class is shown through two studies, based on the analysis of real datasets, aiming at determining the effect

of prices and promotional activities on sales of canned tuna and at evaluating the link between tourism flows and attendance at museums and monuments. The effectiveness of an approach based on the proposed model class in terms of parameter recovery and classification recovery is demonstrated through Monte Carlo studies. The paper is organised as follows. Section 2.1 defines the novel class of 72 cluster-weighted models. Section 2.2 shows how the models belonging to this class relate to some existing models. Information on model identifiability is provided in Section 2.3. Details about the ML estimation are given in Section 2.4 75 and the Appendices. The initialisation and convergence of the ECM algorithm and the issue of model selection are treated in Sections 2.5 and 2.6. Parsimonious 77 models are introduced in Section 2.7. Results of the analyses of simulated and real datasets are summarised in Sections 3 and 4, respectively. Section 5 provides some concluding remarks.

81 2. Multivariate seemingly unrelated linear cluster-weighted analysis

2.1. Multivariate seemingly unrelated linear cluster-weighted models

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Following Dang et al. (2017), in a cluster-weighted model the random vectors \mathbf{X}_i and \mathbf{Y}_i containing the P covariates and the D responses for the ith observation, respectively, come from a population Ω which is assumed to be partitioned into K disjoint groups $\Omega_1, \ldots, \Omega_K$. Thus, $\Omega = \Omega_1 \cup \ldots \cup \Omega_K$; furthermore, $\Omega_k \cap \Omega_{k'} = \emptyset \ \forall k \neq k'$. In the models proposed here both \mathbf{X}_i and \mathbf{Y}_i are continuous random vectors, \mathbf{X}_i takes values in \mathbb{R}^P , \mathbf{Y}_i takes values in \mathbb{R}^D and the probability density function (p.d.f.) of $(\mathbf{X}_i, \mathbf{Y}_i)$ can be written as

$$f(\mathbf{x}_i, \mathbf{y}_i) = \sum_{k=1}^{K} \pi_k f(\mathbf{x}_i | \Omega_k) f(\mathbf{y}_i | \mathbf{x}_i, \Omega_k),$$
(1)

where $\pi_k = \mathbb{P}(\Omega_k)$ is the mixing weight and represents the prior probability of the kth group, $f(\mathbf{x}_i|\Omega_k)$ is the p.d.f. of \mathbf{X}_i given Ω_k and $f(\mathbf{y}_i|\mathbf{x}_i,\Omega_k)$ is the conditional p.d.f. of the response \mathbf{Y}_i given the value \mathbf{x}_i of the covariates \mathbf{X}_i and the group Ω_k . As far as the mixing weights are concerned, they are supposed to

be positive $(\pi_k > 0 \ \forall k)$; in addition, they have to sum to 1 $(\sum_{k=1}^K \pi_k = 1)$. Here $\mathbf{X}_i | \Omega_k$ is assumed to follow a P-variate normal distribution with mean vector $\mu_{\mathbf{X}_k}$ and covariance matrix $\Sigma_{\mathbf{X}_k}$, $k=1,\ldots,K$. Thus, the expected values, variances and covariances of $\mathbf{X}_i | \Omega_k$ are equal for all observations coming from group Ω_k , while they are different for observations belonging to other groups. 99 As far as $\mathbf{Y}_i|(\mathbf{X}_i=\mathbf{x}_i,\Omega_k)$ is concerned, its distribution is modelled using a 100 D-variate normal distribution with conditional expected vector given by some 101 linear transformation of \mathbf{x}_i and covariance matrix $\Sigma_{\mathbf{Y}_k}$. Thus, variances and 102 covariances of $\mathbf{Y}_i|(\mathbf{X}_i=\mathbf{x}_i,\Omega_k)$ are equal for observations coming from the same 103 group; the expected values of $\mathbf{Y}_i|(\mathbf{X}_i=\mathbf{x}_i,\Omega_k)$ for such observations vary with 104 the observations. Furthermore, different correlation structures among both the 105 covariates and the responses across the K groups are assumed. 106

In order to describe how a cluster-weighted model with a different vector of covariates for each response can be obtained, the following additional no-108 tation is required. Suppose that only P_d of the P covariates $(P_d \leq P)$ are 109 considered to be relevant for the prediction of the dth response. Thus, let 110 $\mathbf{x}_{id} = (x_{i,d_1}, x_{i,d_2}, \dots, x_{i,d_{P_d}})'$ be the vector composed of the values of such 111 P_d covariates for the *i*th observation and $\mathbf{x}_{id}^* = (1, \mathbf{x}_{id}')'$. Furthermore, let $\boldsymbol{\beta}_{kd} = (\beta_{kd_1}, \beta_{kd_2}, \dots, \beta_{kd_{P_d}})'$ be the P_d -dimensional vector of regression coefficients capturing the linear effect of these P_d covariates on the dth response in the kth group, and $\beta_{kd}^* = (\beta_{kd0}, \beta'_{kd})'$. Then, the vector containing all linear effects on the D responses in the kth group is given by $\boldsymbol{\beta}_k^* = (\boldsymbol{\beta}_{k1}^{*\prime}, \dots, \boldsymbol{\beta}_{kd}^{*\prime}, \dots, \boldsymbol{\beta}_{kD}^{*\prime})';$ the length of this vector is (P^*+D) , where $P^*=\sum_{d=1}^D P_d$. Finally, the following $(P^* + D) \times D$ partitioned matrix is required:

$$\mathcal{X}_i = egin{bmatrix} \mathbf{x}_{i1}^* & \mathbf{0}_{P_1+1} & \dots & \mathbf{0}_{P_1+1} \ \mathbf{0}_{P_2+1} & \mathbf{x}_{i2}^* & \dots & \mathbf{0}_{P_2+1} \ dots & dots & dots \ \mathbf{0}_{P_D+1} & \mathbf{0}_{P_D+1} & \dots & \mathbf{x}_{iD}^* \end{bmatrix},$$

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with $\mathbf{0}_{P_d+1}$ denoting the (P_d+1) -dimensional null vector. With this notation,

the conditional expected vector of $\mathbf{Y}_i|(\mathbf{X}_i=\mathbf{x}_i,\Omega_k)$ is given by

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$$\boldsymbol{\mu}_{\mathbf{Y}_{k}}(\mathbf{x}_{i};\boldsymbol{\beta}_{k}^{*}) = \mathcal{X}_{i}^{\prime}\boldsymbol{\beta}_{k}^{*} = (\mathbf{x}_{i1}^{*\prime}\boldsymbol{\beta}_{k1}^{*}, \dots, \mathbf{x}_{id}^{*\prime}\boldsymbol{\beta}_{kd}^{*}, \dots, \mathbf{x}_{iD}^{*\prime}\boldsymbol{\beta}_{kD}^{*})^{\prime}. \tag{2}$$

According to this equation, the conditional expected value of the dth response within the kth group is given by the linear term $\mathbf{x}_{id}^{*\prime}\boldsymbol{\beta}_{kd}^{*}$, which only depends on the P_d covariates included in the vector \mathbf{x}_{id} . It is worth noting that the regression coefficients vary across groups, which means that the effect of the covariates on the responses changes with the groups. Embedding all these assumptions into model (1) leads to

$$f(\mathbf{x}_{i}, \mathbf{y}_{i}; \boldsymbol{\psi}) = \sum_{k=1}^{K} \pi_{k} \phi_{P} \left(\mathbf{x}_{i}; \boldsymbol{\mu}_{\mathbf{X}_{k}}, \boldsymbol{\Sigma}_{\mathbf{X}_{k}} \right) \phi_{D} \left(\mathbf{y}_{i} | \mathbf{x}_{i}; \boldsymbol{\mathcal{X}}_{i}' \boldsymbol{\beta}_{k}^{*}, \boldsymbol{\Sigma}_{\mathbf{Y}_{k}} \right),$$
(3)

where ϕ_P (ϕ_D) represents the p.d.f. of a P-variate (D-variate) Gaussian random vector, $\boldsymbol{\psi} = \{\pi_1, \dots, \pi_K, \boldsymbol{\mu}_{\mathbf{X}_1}, \dots, \boldsymbol{\mu}_{\mathbf{X}_K}, \boldsymbol{\Sigma}_{\mathbf{X}_1}, \dots, \boldsymbol{\Sigma}_{\mathbf{X}_K}, \boldsymbol{\beta}_1^*, \dots, \boldsymbol{\beta}_K^*, \boldsymbol{\Sigma}_{\mathbf{Y}_1}, \dots, \boldsymbol{\Sigma}_{\mathbf{Y}_K} \}$ denotes the set of all model parameters and $\boldsymbol{\Psi}$ is the parameter space.

The number of free parameters in $\boldsymbol{\psi}$ is $K - 1 + K(P + P^* + D) + K[P(P + 1)/2 + D(D+1)/2]$, which is the sum of the unknown mixture weights, expected values, variances and covariances.

It is worth stressing that the model in equation (3) differs from the model proposed by Dang et al. (2017) because of a different definition of the linear term for the conditional expected value of $\mathbf{Y}_i|(\mathbf{X}_i=\mathbf{x}_i,\Omega_k)$. If all the P covariates are considered to be relevant for the prediction of all responses, that is $\mathbf{x}_{id}=\mathbf{x}_i$ $\forall d$, then $\mathbf{x}_{id}^*=\mathbf{x}_i^*$ $\forall d$, where $\mathbf{x}_i^*=(1,\mathbf{x}_i')'$, and the following equality holds:

$$\mathcal{X}_i = \mathbf{I}_D \otimes \mathbf{x}_i^*$$
,

where I_D is the identity matrix of order D and \otimes denotes the Kronecker product operator (see, e.g., Magnus and Neudecker, 1988). Then, equation (2) can be rewritten as

$$\boldsymbol{\mu}_{\mathbf{Y}_i}(\mathbf{x}_i; \boldsymbol{\beta}_k^*) = (\mathbf{I}_D \otimes \mathbf{x}_i^*)' \boldsymbol{\beta}_k^* = \mathbf{B}_k' \mathbf{x}_i^*, \ k = 1, \dots, K,$$

where $\mathbf{B}_k = [\boldsymbol{\beta}_{k1}^* \cdots \boldsymbol{\beta}_{kd}^* \cdots \boldsymbol{\beta}_{kD}^*]$, thus leading to the multivariate Gaussian cluster-weighted model introduced by Dang et al. (2017). As illustrated in Section 1, seemingly unrelated regression models can be considered as multivariate

regression models in which prior information about the absence of certain covariates for the prediction of certain responses is explicitly taken into consideration
(Srivastava and Giles, 1987). Thus, equation (3) can also be seen as a multivariate Gaussian cluster-weighted model in which some regression coefficients are
constrained to be a priori equal to zero. To the best of the authors' knowledge,
the inclusion of such constraints in the multivariate Gaussian cluster-weighted
model framework has not been addressed yet.

2.2. Relationships with linear clusterwise regression models

Under suitable conditions, it is possible to establish some relationships between the multivariate seemingly unrelated Gaussian linear cluster-weighted
models just introduced and some Gaussian linear clusterwise regression models.

In Section 2.1 it has been highlighted that models (3) assume that $\mathbf{X}_i | \Omega_k \sim$ $N_P(\boldsymbol{\mu}_{\mathbf{X}_k}, \boldsymbol{\Sigma}_{\mathbf{X}_k})$, for $k = 1, \dots, K$. If the p.d.f of $\mathbf{X}_i | \Omega_k$ does not depend on group Ω_k , i.e., $\phi_P(\mathbf{x}_i; \boldsymbol{\mu}_{\mathbf{X}_k}, \boldsymbol{\Sigma}_{\mathbf{X}_k}) = \phi_P(\mathbf{x}_i; \boldsymbol{\mu}_{\mathbf{X}}, \boldsymbol{\Sigma}_{\mathbf{X}})$ for every $k = 1, \dots, K$,
then equation (3) can also be written as

$$f(\mathbf{x}_{i}, \mathbf{y}_{i}; \boldsymbol{\psi}) = \phi_{P}(\mathbf{x}_{i}; \boldsymbol{\mu}_{\mathbf{X}}, \boldsymbol{\Sigma}_{\mathbf{X}}) \sum_{k=1}^{K} \pi_{k} \phi_{D}(\mathbf{y}_{i} | \mathbf{x}_{i}; \mathcal{X}_{i}' \boldsymbol{\beta}_{k}^{*}, \boldsymbol{\Sigma}_{\mathbf{Y}_{k}}),$$
(4)

165 where

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$$f(\mathbf{y}_{i}|\mathbf{x}_{i};\tilde{\boldsymbol{\psi}}) = \sum_{k=1}^{K} \pi_{k} \phi_{D}\left(\mathbf{y}_{i}|\mathbf{x}_{i}; \mathcal{X}_{i}' \boldsymbol{\beta}_{k}^{*}, \boldsymbol{\Sigma}_{\mathbf{Y}_{k}}\right),$$
 (5)

with $\tilde{\psi} = \{\pi_1, \dots, \pi_K, \boldsymbol{\beta}_1^*, \dots, \boldsymbol{\beta}_K^*, \boldsymbol{\Sigma}_{\mathbf{Y}_1}, \dots, \boldsymbol{\Sigma}_{\mathbf{Y}_K}\}$, is the seemingly unrelated Gaussian clusterwise linear regression model described in Galimberti and Soffritti (2020). This means that the assignment of the data points to the groups is independent of the covariates; such a condition is also known as assignment independence (see, e.g., Hennig, 2000). Furthermore, if the researcher sets $\mathbf{x}_{id} = \mathbf{x}_i$ $\forall d$ (i.e., all the P covariates are assumed to be relevant for the prediction of all responses), then equation (5) leads to the traditional multivariate Gaussian clusterwise linear regression models (Jones and McLachlan, 1992). Thus, when in equation (3) the following conditions hold true: $\boldsymbol{\mu}_{\mathbf{X}_k} = \boldsymbol{\mu}_{\mathbf{X}}$, $\boldsymbol{\Sigma}_{\mathbf{X}_k} = \boldsymbol{\Sigma}_{\mathbf{X}}$ for $k = 1, \dots, K$, then the information about the K disjoint groups $\Omega_1, \dots, \Omega_K$

that compose the population Ω can be equivalently obtained either from the analysis of the conditional p.d.f. $f(\mathbf{y}_i|\mathbf{x}_i;\tilde{\boldsymbol{\psi}})$ through seemingly unrelated linear clusterwise models or from the analysis of the joint p.d.f. $f(\mathbf{x}_i,\mathbf{y}_i;\boldsymbol{\psi})$ through seemingly unrelated linear cluster-weighted models.

Furthermore, when the following conditions hold true: i) the conditional distribution of $Y_{id}|\mathbf{X}_i = \mathbf{x}_i$ changes with K_d disjoint groups $\Omega_{d1}, \ldots, \Omega_{dk_d}, \ldots, \Omega_{dK_d}$ that compose the population Ω for $d=1,\ldots,D;\ ii$) these D partitions of Ω associated with the D responses are mutually independent (i.e., the population is characterised by D independent cluster structures) (Galimberti and Soffritti, 2007); iii) the assignment independence condition holds true for each of these groupings, then the following model can be defined:

$$f(\mathbf{x}_{i}, \mathbf{y}_{i}; \boldsymbol{\psi}) = \phi_{P}(\mathbf{x}_{i}; \boldsymbol{\mu}_{\mathbf{X}}, \boldsymbol{\Sigma}_{\mathbf{X}}) \prod_{d=1}^{D} \sum_{k_{d}=1}^{K_{d}} \pi_{k_{d}} \phi_{1} \left(y_{id} | \mathbf{x}_{i}; \mathbf{x}_{id}^{*} \boldsymbol{\beta}_{k_{d}d}^{*}, \sigma_{k_{d}d}^{2} \right), \quad (6)$$

where y_{id} is the dth element of \mathbf{y}_i , $\mathbf{x}_{id}^{*}\boldsymbol{\beta}_{k_d d}^{*}$ and $\sigma_{k_d d}^{2}$ are the conditional expected value and the variance of $Y_{id}|\mathbf{X}_i = \mathbf{x}_i$ within the group Ω_{dk_d} , respectively. Thus, under conditions i)-iii), model (6) holds true and the information about the D independent partitions of the sample observations should be obtained from D univariate seemingly unrelated linear clusterwise regression models.

In the light of the relationships just illustrated, it is possible to conclude 194 that multivariate seemingly unrelated linear cluster-weighted models will be 195 more effective than multivariate seemingly unrelated linear clusterwise regres-196 sion models when the assignment independence condition does not hold true. 197 Furthermore, an analysis based on the proposed models should be carried out 198 rather than D separate analyses, based on D univariate seemingly unrelated 199 linear clusterwise regression models, whenever either the condition of D inde-200 pendent cluster structures or the assignment independence condition do not 201 hold for the examined population. 202

2.3. Model identifiability

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Identifiability is essential for parameter estimation and represents a preliminary requirement for the consistency and other asymptotic properties of the

ML estimator. Generally speaking, several types of non-identifiability can affect finite mixture models. A first type is due to invariance to relabeling the 207 components (also known as label-switching). Furthermore, non-identifiability is caused by potential overfitting associated with empty components or equal 209 components (see, e.g., Frühwirth-Schnatter, 2006, p. 15). Thus, identifiability 210 of finite mixture models may be achieved after imposing suitable constraints 211 on the parameter space. As far as multivariate Gaussian cluster-weighted models are concerned, conditions ensuring their identifiability have been defined by Dang et al. (2017). Those conditions can be easily modified in order to hold 214 true also for the seemingly unrelated Gaussian linear cluster-weighted models 215 defined according to equation (3). 216

The constraints to be imposed on the parameters are $\pi_k > 0 \ \forall k$ and $(\beta_k^*, \Sigma_{\mathbf{Y}_k}) \neq (\beta_h^*, \Sigma_{\mathbf{Y}_h})$ for $k \neq h$. These constraints make it possible to avoid the two types of non-identifiability illustrated above. Thus, in order to ensure identifiability, the following class of seemingly unrelated cluster-weighted models has to be considered:

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$$\mathfrak{F} = \left\{ f(\mathbf{x}, \mathbf{y}; \bar{\boldsymbol{\psi}}) : f(\mathbf{x}, \mathbf{y}; \bar{\boldsymbol{\psi}}) = \sum_{k=1}^{K} \pi_{k} \phi_{P} \left(\mathbf{x}; \boldsymbol{\mu}_{\mathbf{X}_{k}}, \boldsymbol{\Sigma}_{\mathbf{X}_{k}} \right) \phi_{D} \left(\mathbf{y} | \mathbf{x}; \mathcal{X}' \boldsymbol{\beta}_{k}^{*}, \boldsymbol{\Sigma}_{\mathbf{Y}_{k}} \right),$$
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$$(\mathbf{x}, \mathbf{y}) \in \mathbb{R}^{P+D}, \bar{\boldsymbol{\psi}} \in \bar{\boldsymbol{\Psi}}, K \in \mathbb{N} \right\},$$

where $\bar{\Psi}$ is the constrained parameter space, defined as follows:

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$$\bar{\boldsymbol{\Psi}} = \left\{ \bar{\boldsymbol{\psi}} \in \boldsymbol{\Psi} : \pi_k > 0, \sum_{k=1}^K \pi_k = 1, (\boldsymbol{\beta}_k^*, \boldsymbol{\Sigma}_{\boldsymbol{Y}_k}) \neq (\boldsymbol{\beta}_h^*, \boldsymbol{\Sigma}_{\boldsymbol{Y}_h}) \text{ for } k \neq h \right\}.$$

An additional condition for the class \mathfrak{F} to be identifiable is the existence of a set $\mathcal{W} \subseteq \mathbb{R}^P$ having probability equal to one according to the P-dimensional Gaussian distribution such that the following clusterwise regression model

$$\sum_{k=1}^{K} \phi_{M}\left(\mathbf{y}|\mathbf{x}; \mathcal{X}'\boldsymbol{\beta}_{k}^{*}, \boldsymbol{\Sigma}_{\mathbf{Y}_{k}}\right) \alpha_{k}(\mathbf{x}), \ \mathbf{y} \in \mathbb{R}^{D},$$

is identifiable for each fixed $\mathbf{x} \in \mathcal{W}$, where $\alpha_1(\mathbf{x}), ..., \alpha_K(\mathbf{x})$ are positive weights summing to one for each $\mathbf{x} \in \mathcal{W}$. Under this condition, it is possible to prove that the class \mathfrak{F} results to be identifiable in $\mathcal{W} \times \mathbb{R}^D$. The proof of this result can be

easily obtained from the proof of the analogous result for multivariate Gaussian cluster-weighted models (see Dang et al., 2017, Appendix A), by simply changing the linear term for the conditional expected value of $\mathbf{Y}|(\mathbf{X}=\mathbf{x},\Omega_k)$.

2.4. Parameter estimation

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Given a sample $S = \{(\mathbf{x}_1, \mathbf{y}_1), \dots, (\mathbf{x}_I, \mathbf{y}_I)\}$ of I independent observations 237 from model (3), ML estimation of the model parameters ψ can be carried out by means of an ECM algorithm developed under a general framework dealing 239 with incomplete-data problems (Dempster et al., 1977; Meng and Rubin, 1993). 240 The missing information is the specific component of the mixture from which 241 the sample observations come from; such information can be described by the 242 K-dimensional vectors $(\mathbf{z}_1, \dots, \mathbf{z}_I)$, where $\mathbf{z}_i = (z_{i1}, \dots, z_{iK})'$ with $z_{ik} = 1$ if the ith observation comes from the kth component and $z_{ik} = 0$ otherwise, for k = $1, \ldots, K$. Then, the complete data would be $S_c = \{(\mathbf{x}_1, \mathbf{y}_1, \mathbf{z}_1), \ldots, (\mathbf{x}_I, \mathbf{y}_I, \mathbf{z}_I)\}.$ Thus, the likelihood functions derived from the incomplete data and the com-246 plete data are

$$L(\boldsymbol{\psi}|\mathcal{S}) = \prod_{i=1}^{I} \left[\sum_{k=1}^{K} \pi_{k} \phi_{P} \left(\mathbf{x}_{i}; \boldsymbol{\mu}_{\mathbf{X}_{k}}, \boldsymbol{\Sigma}_{\mathbf{X}_{k}} \right) \phi_{D} \left(\mathbf{y}_{i} | \mathbf{x}_{i}; \boldsymbol{\mathcal{X}}_{i}' \boldsymbol{\beta}_{k}^{*}, \boldsymbol{\Sigma}_{\mathbf{Y}_{k}} \right) \right],$$

$$L(\boldsymbol{\psi}|\mathcal{S}_{c}) = \prod_{i=1}^{I} \prod_{k=1}^{K} \left[\pi_{k} \phi_{P} \left(\mathbf{x}_{i}; \boldsymbol{\mu}_{\mathbf{X}_{k}}, \boldsymbol{\Sigma}_{\mathbf{X}_{k}} \right) \phi_{D} \left(\mathbf{y}_{i} | \mathbf{x}_{i}; \boldsymbol{\mathcal{X}}_{i}' \boldsymbol{\beta}_{k}^{*}, \boldsymbol{\Sigma}_{\mathbf{Y}_{k}} \right) \right]^{z_{ik}},$$

$$249 \qquad L(\boldsymbol{\psi}|\mathcal{S}_{c}) = \prod_{i=1}^{I} \prod_{k=1}^{K} \left[\pi_{k} \phi_{P} \left(\mathbf{x}_{i}; \boldsymbol{\mu}_{\mathbf{X}_{k}}, \boldsymbol{\Sigma}_{\mathbf{X}_{k}} \right) \phi_{D} \left(\mathbf{y}_{i} | \mathbf{x}_{i}; \boldsymbol{\mathcal{X}}_{i}' \boldsymbol{\beta}_{k}^{*}, \boldsymbol{\Sigma}_{\mathbf{Y}_{k}} \right) \right]^{z_{ik}},$$

respectively; the complete-data log-likelihood function employed in the ECM algorithm for the computation of the parameter estimates is

$$\ell(\boldsymbol{\psi}|\mathcal{S}_{c}) = \sum_{i=1}^{I} \sum_{k=1}^{K} z_{ik} \Big[\ln \pi_{k} + \ln \phi_{P} \left(\mathbf{x}_{i}; \boldsymbol{\mu}_{\mathbf{X}_{k}}, \boldsymbol{\Sigma}_{\mathbf{X}_{k}} \right) + \ln \phi_{D} \left(\mathbf{y}_{i} | \mathbf{x}_{i}; \boldsymbol{\mathcal{X}}_{i}' \boldsymbol{\beta}_{k}^{*}, \boldsymbol{\Sigma}_{\mathbf{Y}_{k}} \right) \Big].$$
(7)

The hth iteration of the E-step in the ECM algorithm consists in calculating
the conditional expectation of the complete-data log-likelihood (7) on the basis
of the current estimate $\hat{\psi}^{(h)}$ of the model parameters ψ :

$$\mathbb{E}_{\hat{\boldsymbol{\psi}}^{(h)}}\left[\ell(\boldsymbol{\psi}|\mathcal{S}_c)\right] = \sum_{i=1}^{I} \sum_{k=1}^{K} \hat{\tau}_{ik}^{(h)} \left[\ln \hat{\pi}_k^{(h)} + Q_1\left(\boldsymbol{\mu}_{\mathbf{X}_k}, \boldsymbol{\Sigma}_{\mathbf{X}_k} | \hat{\boldsymbol{\psi}}^{(h)}\right)\right] + Q_2\left(\boldsymbol{\beta}_k^*, \boldsymbol{\Sigma}_{\mathbf{Y}_k} | \hat{\boldsymbol{\psi}}^{(h)}\right)\right], \tag{8}$$

259 where

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$$Q_{1}\left(\boldsymbol{\mu}_{\mathbf{X}_{k}}, \boldsymbol{\Sigma}_{\mathbf{X}_{k}} | \hat{\boldsymbol{\psi}}^{(h)}\right) = \frac{1}{2} \left[-P \ln(2\pi) - \ln |\hat{\boldsymbol{\Sigma}}_{\mathbf{X}_{k}}^{(h)}| -\left(\mathbf{x}_{i} - \hat{\boldsymbol{\mu}}_{\mathbf{X}_{k}}^{(h)}\right)' \hat{\boldsymbol{\Sigma}}_{\mathbf{X}_{k}}^{(h)(-1)} \left(\mathbf{x}_{i} - \hat{\boldsymbol{\mu}}_{\mathbf{X}_{k}}^{(h)}\right) \right],$$

$$Q_{2}\left(\boldsymbol{\beta}_{k}^{*}, \boldsymbol{\Sigma}_{\mathbf{Y}_{k}} | \hat{\boldsymbol{\psi}}^{(h)}\right) = \frac{1}{2} \left[-D \ln(2\pi) - \ln |\hat{\boldsymbol{\Sigma}}_{\mathbf{Y}_{k}}^{(h)}| -\left(\mathbf{y}_{i} - \boldsymbol{\mathcal{X}}_{i}' \hat{\boldsymbol{\beta}}_{k}^{*(h)}\right)' \hat{\boldsymbol{\Sigma}}_{\mathbf{Y}_{k}}^{(h)(-1)} \left(\mathbf{y}_{i} - \boldsymbol{\mathcal{X}}_{i}' \hat{\boldsymbol{\beta}}_{k}^{*(h)}\right) \right],$$

$$= -\left(\mathbf{y}_{i} - \boldsymbol{\mathcal{X}}_{i}' \hat{\boldsymbol{\beta}}_{k}^{*(h)}\right)' \hat{\boldsymbol{\Sigma}}_{\mathbf{Y}_{k}}^{(h)(-1)} \left(\mathbf{y}_{i} - \boldsymbol{\mathcal{X}}_{i}' \hat{\boldsymbol{\beta}}_{k}^{*(h)}\right) \right],$$

and $\hat{\tau}_{ik}^{(h)}$ provides the posterior probability (evaluated using the current estimate $\hat{\psi}^{(h)}$) that $(\mathbf{x}_i, \mathbf{y}_i)$ is generated from the kth component of the mixture, that is

$$\hat{\tau}_{ik}^{(h)} = \mathbb{E}_{\hat{\boldsymbol{\psi}}^{(h)}} \left[Z_{ik} | \mathbf{x}_{i}, \mathbf{y}_{i} \right] = \mathbb{P}_{\hat{\boldsymbol{\psi}}^{(h)}} \left\{ Z_{ik} = 1 | \mathbf{x}_{i}, \mathbf{y}_{i} \right\}$$

$$= \frac{\hat{\pi}_{k}^{(h)} \phi_{P} \left(\mathbf{x}_{i}; \hat{\boldsymbol{\mu}}_{\mathbf{X}_{k}}^{(h)}, \hat{\boldsymbol{\Sigma}}_{\mathbf{X}_{k}}^{(h)} \right) \phi_{D} \left(\mathbf{y}_{i} | \mathbf{x}_{i}; \mathcal{X}_{i}' \hat{\boldsymbol{\beta}}_{k}^{*(h)}, \hat{\boldsymbol{\Sigma}}_{\mathbf{Y}_{k}}^{(h)} \right)}{\sum_{k'=1}^{K} \hat{\pi}_{k'}^{(h)} \phi_{P} \left(\mathbf{x}_{i}; \hat{\boldsymbol{\mu}}_{\mathbf{X}_{k'}}^{(h)}, \hat{\boldsymbol{\Sigma}}_{\mathbf{X}_{k'}}^{(h)} \right) \phi_{D} \left(\mathbf{y}_{i} | \mathbf{x}_{i}; \mathcal{X}_{i}' \hat{\boldsymbol{\beta}}_{k'}^{*(h)}, \hat{\boldsymbol{\Sigma}}_{\mathbf{Y}_{k'}}^{(h)} \right)}. \tag{9}$$

The (h+1)th update of $\hat{\boldsymbol{\psi}}^{(h)}$ is obtained by a sequence of CM-steps involved in the ECM algorithm. These steps are meant to maximise the conditional expectation of $\ell(\boldsymbol{\psi}|\mathcal{S}_c)$ with respect to $\boldsymbol{\psi}$. This maximisation can be achieved by setting the first order derivatives of $\mathbb{E}\left[\ell(\boldsymbol{\psi}|\mathcal{S}_c)\right]$ equal to zero and then solving the resulting system of equations with respect to the parameters of interest. Since this expected value can be decomposed in a sum of three terms, each one depending on a specific set of parameters (see equation (8)), maximisation can be carried out separately for each set of parameters. The resulting updates of $\hat{\pi}_k^{(h)}, \hat{\mu}_{\mathbf{X}_k}^{(h)}, \hat{\Sigma}_{\mathbf{X}_k}^{(h)}, k = 1, \dots, K$ are:

$$\hat{\pi}_k^{(h+1)} = \frac{1}{I} \sum_{i=1}^{I} \hat{\tau}_{ik}^{(h)}, \tag{10}$$

$$\hat{\boldsymbol{\mu}}_{\mathbf{X}_{k}}^{(h+1)} = \frac{\sum_{i=1}^{I} \hat{\tau}_{ik}^{(h)} \mathbf{x}_{i}}{\sum_{i=1}^{I} \hat{\tau}_{ik}^{(h)}}, \tag{11}$$

$$\hat{\Sigma}_{\mathbf{X}_{k}}^{(h+1)} = \frac{\sum_{i=1}^{I} \hat{\tau}_{ik}^{(h)} \left(\mathbf{x}_{i} - \hat{\boldsymbol{\mu}}_{\mathbf{X}_{k}}^{(h+1)}\right) \left(\mathbf{x}_{i} - \hat{\boldsymbol{\mu}}_{\mathbf{X}_{k}}^{(h+1)}\right)'}{\sum_{i=1}^{I} \hat{\tau}_{ik}^{(h)}}.$$
 (12)

Such updates coincide with the ones reported in Dang et al. (2017). The CM-

steps to update the remaining parameters are (see Appendix A for a proof)

$$\hat{\boldsymbol{\beta}}_{k}^{*(h+1)} = \left[\sum_{i=1}^{I} \hat{\tau}_{ik}^{(h)} \mathcal{X}_{i} \left(\hat{\boldsymbol{\Sigma}}_{\mathbf{Y}_{k}}^{(h)} \right)^{-1} \mathcal{X}_{i}' \right]^{-1} \left[\sum_{i=1}^{I} \hat{\tau}_{ik}^{(h)} \mathcal{X}_{i} \left(\hat{\boldsymbol{\Sigma}}_{\mathbf{Y}_{k}}^{(h)} \right)^{-1} \mathbf{y}_{i} \right], (13)$$

$$\hat{\Sigma}_{\mathbf{Y}_{k}}^{(h+1)} = \frac{\sum_{i=1}^{I} \hat{\tau}_{ik}^{(h)} \left(\mathbf{y}_{i} - \mathcal{X}_{i}' \hat{\boldsymbol{\beta}}_{k}^{*(h+1)}\right) \left(\mathbf{y}_{i} - \mathcal{X}_{i}' \hat{\boldsymbol{\beta}}_{k}^{*(h+1)}\right)'}{\sum_{i=1}^{I} \hat{\tau}_{ik}^{(h)}}.$$
 (14)

It is worth noting that the matrix $\sum_{i=1}^{I} \hat{\tau}_{ik}^{(h)} \mathcal{X}_i \left(\hat{\Sigma}_{\mathbf{Y}_k}^{(h)}\right)^{-1} \mathcal{X}_i'$ has to be nonsingular in order for the update $\hat{\beta}_k^{*(h+1)}$ in equation (13) to exist. In addition, Appendix B shows that equation (13) is equivalent to the expression reported 287 in Dang et al. (2017) for the updates of the regression coefficient matrix when 288 $\mathbf{x}_{id} = \mathbf{x}_i \ \forall d$. As a consequence, in this special case the ECM algorithm de-289 scribed in this section reduces to the EM algorithm described in Dang et al. (2017). Finally, once the convergence is reached, the ECM algorithm also provides estimates of the posterior probabilities according to equation (9), which 292 can be used to partition the I observations into K clusters, by assigning each 293 observation to the component showing the highest posterior probability. 294

Difficulties with this ECM algorithm can arise when matrices $\hat{m{\Sigma}}_{{f X}_k}^{(h+1)}$ and $\hat{\Sigma}_{\mathbf{Y}_k}^{(h+1)}$ in equations (12) and (14) are singular or nearly singular. Another difficulty with ML estimation of Gaussian mixture models is the unboundedness 297 of the likelihood function (see, e.g. Frühwirth-Schnatter, 2006, p. 173). A way 298 to deal with these problems is to introduce suitable constraints on the param-299 eter space Ψ and to perform the estimation under a constrained Ψ (see, e.g. Ingrassia and Rocci, 2011; Rocci et al., 2018). All the analyses illustrated in this paper have been carried out through an implementation of the proposed 302 ECM algorithm, which also allows the estimation of the multivariate linear 303 cluster-weighted models introduced by Dang et al. (2017), in the R environment. 304 Such an implementation embeds suitable constraints on the eigenvalues of both $\hat{\Sigma}_{\mathbf{X}_k}^{(h+1)}$ and $\hat{\Sigma}_{\mathbf{Y}_k}^{(h+1)}$ for $k=1,\ldots,K$. Namely, following Dang et al. (2017), all estimated covariance matrices have been required to have eigenvalues greater than 307 the conservative bound 10^{-20} ; furthermore, the ratio between the smallest and 308 the largest eigenvalues of such matrices is required to be not lower than 10^{-10} .

Finally, in order to avoid problems associated with the invariance of a mixture distribution to relabeling its components (see, e.g., Frühwirth-Schnatter, 2006, p. 15), the K estimated components of the model (3) have been labeled according to the estimated prior probabilities taken in non-decreasing order.

2.5. Initialisation and convergence of the ECM algorithm

A crucial point of any ECM algorithm is the choice of the starting values 315 for the model parameters (i.e., $\hat{\psi}^{(0)}$). An approach based on multiple random 316 initialisations and multiple executions of the ECM algorithm could be adopted. 317 Approaches based on non-random choices can be employed. A solution could 318 be obtained by resorting to the following two-step strategy. In the first step 319 a mixture of K Gaussian models is estimated for the joint distribution all covariates and responses. This task can be carried out, for example, by resorting 32 to the mclust package (Scrucca et al., 2017) for the R environment. The K 322 prior probabilities, mean vectors and covariance matrices for the predictors es-323 timated in this way are used as $\hat{\pi}_k^{(0)}$, $\hat{\mu}_{\mathbf{X}_k}^{(0)}$ and $\hat{\Sigma}_{\mathbf{X}_k}^{(0)}$, for $k = 1, \dots, K$. In the second step $\hat{\boldsymbol{\beta}}_k^{*(0)}$ and $\hat{\boldsymbol{\Sigma}}_{\mathbf{Y}_k}^{(0)}$ are obtained from an estimate of the parameters of the conditional distribution of the responses given the predictors based on the 326 fitting of a seemingly unrelated Gaussian linear regression model to the sam-327 ple observations that have been assigned to the kth component of the mixture 328 model estimated in the first step. The R package systemfit (Henningsen and Hamann, 2007) can be exploited to perform this task. Another way to obtain $\hat{\pi}_k^{(0)}, \hat{\mu}_{\mathbf{X}_k}^{(0)}$ and $\hat{\Sigma}_{\mathbf{X}_k}^{(0)}$, for $k=1,\ldots,K$ could be based on the fitting of a mixture 33 of K Gaussian models for the marginal distribution of the covariates in the first 332 step of the previous strategy while keeping the second step unchanged. In all 333 analyses reported in this paper involving either models defined in equation (3) 334 or models introduced by Dang et al. (2017), both these strategies have been simultaneously employed; thus, two different initialisations have been considered 336 for each analysed dataset. Then, the ECM algorithm has been initialised with 337 the strategy leading to the largest value of the incomplete log-likelihood. 338

In the R function employed for the parameter estimation in all analyses

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summarised in this paper, the convergence of the ECM algorithm has been 340 evaluated through a criterion based on the Aitken acceleration (Aitken, 1926) 343 which consists in stopping the ECM algorithm when $\left|l_A^{(h+1)} - \ell(\hat{\psi}^{(h)}|\mathcal{S})\right| < \epsilon$, where $l_A^{(h+1)}$ is the (h+1)th Aitken accelerated estimate of the log-likelihood limit and $\ell(\hat{\psi}^{(h)}|\mathcal{S})$ is the value of the incomplete log-likelihood at the hth 344 iteration (see, e.g., McNicholas, 2010, for more details). Such criterion can avoid 345 premature stops associated with the use of lack of progress stopping criteria, such as the one based on the difference between the log-likelihood values at two consecutive steps of the ECM algorithm. The maximum number of iterations 348 for the ECM algorithm and the value for ϵ have been set equal to 500 and 10^{-8} , 349 respectively. 350

351 2.6. Model selection

The ECM algorithm described in Section 2.4 performs the ML estimation 352 for a given value of K. However, in most practical applications, the number 353 of groups is not known and must be determined from the data \mathcal{S} . A common solution to this task is obtained by resorting to model selection criteria which 355 allows to trade-off the fit (measured by $l_M(\hat{\psi}|\mathcal{S})$, the maximum of the incomplete 356 loglikelihood of model M) and complexity (given by $npar_M$, the number of 357 free parameters in model M) (see, e.g., Frühwirth-Schnatter, 2006, subsections 4.4.2-4.4.3). In the context of Gaussian mixture models and Gaussian cluster-359 weighted models (see, e.g., Fraley and Raftery, 2002; Dang et al., 2017), the 360 Bayesian Information Criterion (BIC) (Schwarz, 1978) has performed well and 361 is commonly employed. It can be computed as follows: $BIC_M = -2l_M(\hat{\psi}|\mathcal{S}) +$ 362 $npar_M \ln I$. Given a collection of competing fitted candidate models, the one that 363 minimises BIC_M is preferred. Model selection criteria that also consider the quality of the estimated partition of the sample observations represent another possible solution (see, e.g., Frühwirth-Schnatter, 2006, subsection 7.1.4). 366

2.7. Parsimonious models

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As the number of free parameters in equation (3) incresses quadratically with both the number of responses and the number of predictors, analyses

based on the proposed models can become unfeasible in practical applications. This problem can be overcome by introducing constraints on the elements of 371 the covariance matrices $\Sigma_{\mathbf{X}_k}$ and $\Sigma_{\mathbf{Y}_k}$ $(k=1,\ldots,K)$ according to the ap-372 proach illustrated in Celeux and Govaert (1995). In this approach, the follow-373 ing eigen-decomposition of the covariance matrix $\Sigma_{\mathbf{X}_k}$ has to be considered: 374 $\mathbf{\Sigma}_{\mathbf{X}_k} = \alpha_k \mathbf{D}_k \mathbf{A}_k \mathbf{D}_k'$, where $\alpha_k = |\mathbf{\Sigma}_{\mathbf{X}_k}|^{1/D}$, \mathbf{A}_k is the diagonal matrix con-375 taining the eigenvalues of $\Sigma_{\mathbf{X}_k}$ (normalised in such a way that $|\mathbf{A}_k|=1$) and \mathbf{D}_k is the matrix of eigenvectors of $\Sigma_{\mathbf{X}_k}$. Thus, variances and covariances in $\Sigma_{\mathbf{X}_k}$ can be obtained from α_k , \mathbf{A}_k and \mathbf{D}_k . From a geometrical point of view, 378 such parameters determine the volume, shape and orientation of the kth cluster 379 of observations with respect to the predictors. By constraining one or more 380 of these three parameters to be equal across components, 14 different covari-38: ance structures for the predictors in models (3) with K > 1 can be determined (see Celeux and Govaert, 1995, for more details). Additional information about 383 these parameterisations can be found in Table 1. The application of the same 384 approach to the covariance matrices $\Sigma_{\mathbf{Y}_k}$, $k = 1, \ldots, K$ leads to a class of 196 385 different models for any given K > 1. Equations (12) and (14) represent the 386 solutions for the model in which the covariance structures of both predictors 387 and responses are fully unconstrained. For all other parsimonious models, the 388 CM-step updates for the estimation of $\Sigma_{\mathbf{X}_k}$ and $\Sigma_{\mathbf{Y}_k}$ in the ECM algorithm 389 have to be modified; these modified updates can be computed either in closed 390 form or using iterative procedures, depending on the specific parameterisation to be employed (see Celeux and Govaert, 1995, for more details). The CM-step updates $\hat{\Sigma}_{\mathbf{X}_k}^{(h+1)}$ and $\hat{\Sigma}_{\mathbf{Y}_k}^{(h+1)}$ associated with the parameterisations EVE and VVE 393 can be computed using the F-G algorithm (Flury and Gautschi, 1986) or one of 394 its variants (see, e.g., Lin, 2014). Algorithms which are computationally feasible 395 also in high-dimensional situations have been recently introduced (Browne and 396 McNicholas, 2014a,b). All the experimental results illustrated here and concerning the EVE and VVE parameterisations have been obtained using the algorithms 398 given in Browne and McNicholas (2014a). When K=1, only three covariance 399 structures for both responses and covariates are possible: diagonal with different

Table 1: Parsimonious parameterisations for the component-covariance matrices

Acronym	Model	Distribution	Volume	Shape	Orientation
EEE	$\alpha \mathbf{D} \mathbf{A} \mathbf{D}'$	Ellipsoidal	Equal	Equal	Equal
VVV	$\alpha_k \mathbf{D}_k \mathbf{A}_k \mathbf{D}_k'$	Ellipsoidal	Variable	Variable	Variable
EII	$lpha {f I}$	Spherical	Equal	Equal	_
VII	$\alpha_k \mathbf{I}$	Spherical	Variable	Equal	_
EEI	$\alpha \mathbf{A}$	Diagonal	Equal	Equal	_
VEI	$\alpha_k \mathbf{A}$	Diagonal	Variable	Equal	_
EVI	$\alpha \mathbf{A}_k$	Diagonal	Equal	Variable	_
VVI	$\alpha_k \mathbf{A}_k$	Diagonal	Variable	Variable	_
EEV	$\alpha \mathbf{D}_k \mathbf{A} \mathbf{D}_k'$	Ellipsoidal	Equal	Equal	Variable
VEV	$\alpha_k \mathbf{D}_k \mathbf{A} \mathbf{D}_k'$	Ellipsoidal	Variable	Equal	Variable
EVE	$\alpha \mathbf{D} \mathbf{A}_k \mathbf{D}'$	Ellipsoidal	Equal	Variable	Equal
VVE	$\alpha_k \mathbf{D} \mathbf{A}_k \mathbf{D}'$	Ellipsoidal	Variable	Variable	Equal
VEE	$\alpha_k \mathbf{D} \mathbf{A} \mathbf{D}'$	Ellipsoidal	Variable	Equal	Equal
EVV	$\alpha \mathbf{D}_k \mathbf{A}_k \mathbf{D}_k'$	Ellipsoidal	Equal	Variable	Variable

entries (VI), diagonal with the same entries (EI) and fully unconstrained (VV).

Thus, nine differentially parameterised one-component cluster-weighted models

403 can be obtained.

3. Results from Monte Carlo studies

The main purpose of the studies based on simulated datasets illustrated here is to obtain an evaluation of the effectiveness of the proposed methodology in comparison with the approach introduced by Dang et al. (2017), where the same vector of covariates has to be employed for all responses. Thus, cluster-weighted models belonging to two different classes have been fitted to each simulated dataset: i) models in which all the D responses are assumed to depend on all the P examined covariates (i.e., the models proposed by Dang et al. (2017)); ii) models defined according to equation (3) in which each response has its

specific predictors. From now on, such models have been denoted as CW and 413 SuCW, respectively. A hundred datasets of I = 450 independent observations 414 have been randomly generated from model (3) with D=2 responses, P=3415 predictors and K = 3 components in which the elements of the conditional 416 expected vector (2) are defined as follows: 417

$$E(Y_{i1}|\mathbf{X}_i = \mathbf{x}_i, \Omega_k) = \beta_{k10} + \beta_{k11}x_{i1} + \beta_{k12}x_{i2}, \tag{15}$$

$$E(Y_{i2}|\mathbf{X}_i = \mathbf{x}_i, \Omega_k) = \beta_{k20} + \beta_{k21}x_{i1} + \beta_{k22}x_{i3}.$$
 (16)

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and $\epsilon = 0.375$, respectively.

Thus, the model employed to generate the datasets assumes that the first re-420 sponse Y_1 depends on X_1 and X_2 while Y_2 depends on X_1 and X_3 . Further-421 more, the component-covariance structures of both the predictors and the re-422 sponses are defined using the VVV parameterisation. The specific values of the 423 parameters for the data-generating model are: $\pi_1 = 0.4$, $\pi_2 = 0.35$, $\pi_3 = 0.25$, 424 $\mu_{\mathbf{X}_1} = (0,0,0)', \; \mu_{\mathbf{X}_2} = (2,4,-2)', \; \mu_{\mathbf{X}_3} = \mu_{\mathbf{X}_2} + 2\epsilon \cdot \mathbf{1}_P, \; \text{where } \mathbf{1}_P \; \text{is the}$ $P \times 1$ vector having each element equal to 1, $\boldsymbol{\beta}_1^* = (-2, 0.75, 1, 1, 0.5)$ $\boldsymbol{\beta}_{2}^{*} = (0.5, 1.75, 0.25, 1, 1, 1)', \boldsymbol{\beta}_{3}^{*} = \boldsymbol{\beta}_{2}^{*} + \epsilon \cdot \mathbf{1}_{6}, \boldsymbol{\Sigma}_{\mathbf{X}_{1}} = \begin{pmatrix} 1.72 & -0.18 & 0.27 \\ -0.18 & 1.89 & 0.27 \\ 0.27 & 0.27 & 2.89 \end{pmatrix},$ $\Sigma_{\mathbf{X}_{2}} = \begin{pmatrix} 2.33 & -0.52 & -0.06 \\ -0.52 & 0.88 & -0.34 \\ -0.06 & -0.34 & 1.04 \end{pmatrix}, \ \Sigma_{\mathbf{X}_{3}} = \Sigma_{\mathbf{X}_{2}}, \ \Sigma_{\mathbf{Y}_{1}} = \begin{pmatrix} 1.34 & 0.47 \\ 0.47 & 1.66 \end{pmatrix}, \ \Sigma_{\mathbf{Y}_{2}} = \begin{pmatrix} 0.50 & 0.04 \\ 0.04 & 1.50 \end{pmatrix}, \ \Sigma_{\mathbf{Y}_{3}} = \Sigma_{\mathbf{Y}_{2}}.$ Since the second and third components of the data-generating model only differ in the values of intercepts and regression coefficients and the expected values of the regressors, the separation between such 431 components depends on ϵ . The simulated datasets have been generated using 432

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the following values of ϵ : 0.275, 0.3, 0.325, 0.350 and 0.375; this allows an evalu-

ation of the performances of the approaches based on SuCW and CW models under

different experimental levels of separation between those components. Figures 1

and 2 show the scatterplots for two simulated datasets obtained with $\epsilon = 0.3$

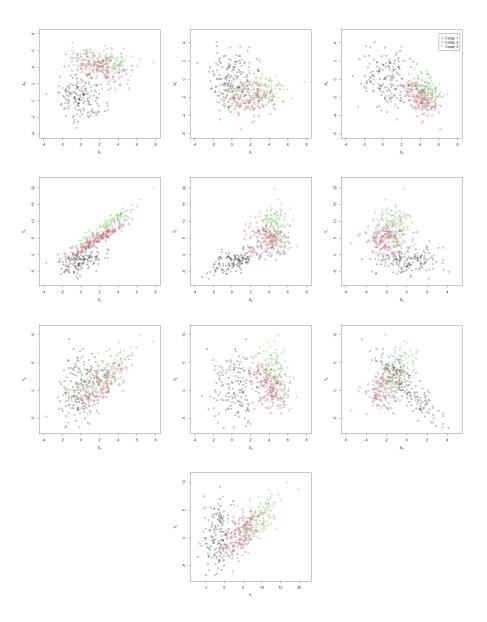


Figure 1: Bivariate scatter plots for pairs of variables in a simulated dataset, $\epsilon=0.3.$

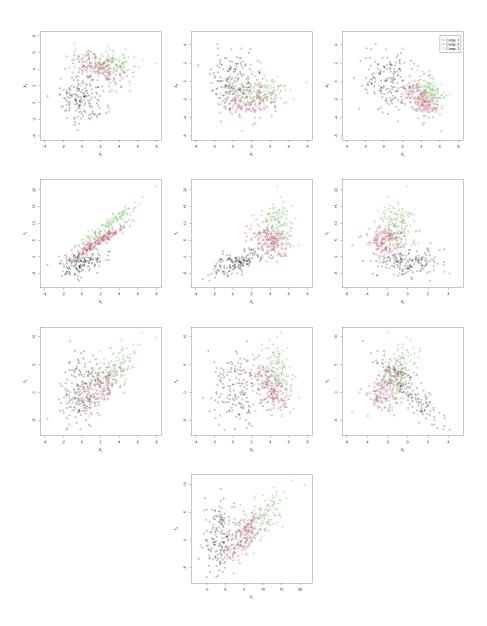


Figure 2: Bivariate scatter plots for pairs of variables in a simulated dataset, $\epsilon=0.375.$

A first analysis has been carried out where the 196 SuCW and CW models with 438 K=3 components associated with all the parameterisations for the component-439 covariance structures of both the predictors and the responses have been fitted to each dataset. It is worth noting that using CW models leads to non-parsimonious 441 specifications for such datasets, as six regression coefficients (two for each com-442 ponent) have been estimated although in fact they are equal to zero. The 443 analysis has been run on an IBM x3750 M4 server with 4 Intel Xeon E5-4620 processors with 8 cores and 128GB RAM. The average execution times (over 100 datasets) for SuCW models have ranged between 2.698 and 35.309 seconds, 446 depending on the specific combination of parameterisations for the component 447 covariance matrices and the value of ϵ . Concerning CW models, the minimum 448 and maximum average execution times have resulted to be equal to 3.382 and 40.710 seconds, respectively. Since the implementation of the ECM algorithm has not been carried out with the goal of being efficient from a computational 451 point of view, these CPU times are merely illustrative and can be reduced using 452 more efficient implementations. For all the models fitted to any dataset, the 453 value of BIC has been computed and the models with the lowest BIC within 454 the two collections of fitted models have been selected. The 100 pairs of models 455 selected as just illustrated, one for each simulated dataset, have been employed 456 to compare the effectiveness of the two approaches. As expected, SuCW models 457 have resulted to be preferable to CW ones. For each dataset $BIC_{\mathtt{SuCW}} < BIC_{\mathtt{CW}}$ for 458 all the examined values of ϵ with the exception of two datasets when $\epsilon = 0.350$. 459 A further evaluation of the two approaches has been performed by examining 460 their ability to recover the true values of the unknown parameters (i.e., param-461 eter recovery). In particular, the attention has been focused on the bias and 462 the root mean squared error (RMSE) for the regression coefficients in equations 463

464 (15) and (16). Namely, the following quantities have been computed

$$Bias\left(\hat{\beta}_{kdp}\right) = \left|\beta_{kdp} - \frac{\sum_{r=1}^{100} \hat{\beta}_{kdp}^{(r)}}{100}\right|, \ k = 1, 2, 3, \ d = 1, 2, \ p = 1, 2,$$

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$$RMSE\left(\hat{\beta}_{kdp}\right) = \sqrt{\frac{\sum_{r=1}^{100} \left(\beta_{kdp} - \hat{\beta}_{kdp}^{(r)}\right)^2}{100}}, \ k = 1, 2, 3, \ d = 1, 2, \ p = 1, 2,$$

where $\hat{\beta}_{kdp}^{(r)}$ is the ML estimate of β_{kdp} obtained from the rth dataset (r = 1,..., 100). Note that CW models contain additional regression coefficients as-468 sociated with the equation-specific irrelevant regressors. The bias and RMSE 469 have been computed also for these additional coefficients, using 0 as their true 470 value. Tables 2 and 3 report the values of bias and RMSE, respectively, ob-471 tained for each value of ϵ . Overall, both approaches tend to provide acceptable 472 results in terms of recovering the true values of the regression coefficients. This 473 is evident for the parameters of the first component. As far as the second and 474 third components are concerned, there seems to be a tendency for SuCW models to perform slightly better than CW models, especially considering the RMSE for 476 low values of ϵ . It is also worth noting that CW models appear to be capable of 477 recognising the presence of irrelevant regressors, as the corresponding estimated 478 regression coefficients are on average very close to 0. However, the RMSE of 479 some of these estimates tend to be large, suggesting a a low precision in the 480 estimation of the effect of some irrelevant regressors. This precision seems to 481 improve as the separation among components increases. 482

The performance of the two approaches has also been evaluated by their abil-483 ity to properly estimate the true classification of the sample observations (i.e., 484 classification recovery). This task has been carried out by means of the adjusted 485 Rand index (ARI) (Hubert and Arabie, 1985). Some summary statistics of this 486 index (over the 100 datasets) for both approaches by the five examined levels 487 of separation are reported in Table 4. These results show that the classification 488 recovery associated with the use of both approaches increases with the level of 489 separation between the second and third components (see the mean and me-490 dian values of ARI in Table 4); on the contrary, the interquartile range and 491

Table 2: Bias for the regression coefficients under SuCW and CW models in the first study.

	$\epsilon = 0$	0.275	$\epsilon =$	0.3	$\epsilon = 0$	0.325	$\epsilon = 0$	0.350	$\epsilon = 0$	0.375
	SuCW	CW	SuCW	CW	SuCW	CW	SuCW	CW	SuCW	CW
$\beta_{111} = 0.75$	0.005	0.003	0.005	0.004	0.006	0.005	0.007	0.005	0.006	0.005
$\beta_{112} = 1$	0.003	0.001	0.002	0.003	0.003	0.003	0.002	0.005	0.002	0.003
$\beta_{121} = 0.5$	0.010	0.013	0.005	0.007	0.005	0.008	0.002	0.006	0.005	0.008
$\beta_{122} = -2$	0.000	0.005	0.002	0.004	0.002	0.004	0.002	0.004	0.002	0.004
$\beta_{211} = 1.75$	0.027	0.090	0.011	0.023	0.001	0.010	0.037	0.002	0.003	0.002
$\beta_{212} = 0.25$	0.067	0.160	0.023	0.040	0.009	0.022	0.010	0.023	0.000	0.002
$\beta_{221} = 1$	0.028	0.039	0.019	0.011	0.013	0.009	0.007	0.008	0.009	0.007
$\beta_{222} = 1$	0.004	0.026	0.028	0.042	0.027	0.023	0.027	0.053	0.028	0.028
$\beta_{311} = 1.75 + \epsilon$	0.002	0.005	0.002	0.003	0.001	0.002	0.003	0.003	0.002	0.003
$\beta_{312} = 0.25 + \epsilon$	0.052	0.100	0.014	0.034	0.010	0.016	0.012	0.016	0.004	0.017
$\beta_{321} = 1 + \epsilon$	0.023	0.065	0.022	0.028	0.013	0.012	0.017	0.018	0.009	0.009
$\beta_{322} = 1 + \epsilon$	0.004	0.020	0.013	0.015	0.016	0.007	0.009	0.011	0.020	0.021
Irrelevant regres	sors									
$\beta_{113} = 0$	_	0.007	_	0.003	_	0.003	_	0.001	_	0.003
$\beta_{123} = 0$	_	0.022	_	0.021	_	0.021	_	0.022	_	0.021
$\beta_{213} = 0$	_	0.060	_	0.016	_	0.015	_	0.006	_	0.002
$\beta_{223} = 0$	_	0.001	_	0.011	_	0.005	_	0.004	_	0.003
$\beta_{313} = 0$	_	0.067	_	0.040	_	0.028	_	0.023	_	0.019
$\beta_{323} = 0$	_	0.082	_	0.011	_	0.002	_	0.007	_	0.003

Table 3: RMSE for the regression coefficients under SuCW and CW models in the first study.

	$\epsilon = 0$	0.275	$\epsilon =$	0.3	$\epsilon = 0$	0.325	$\epsilon = 0$	0.350	$\epsilon = 0$	0.375
	SuCW	CW	SuCW	CW	SuCW	CW	SuCW	CW	SuCW	CW
$\beta_{111} = 0.75$	0.058	0.059	0.058	0.059	0.059	0.059	0.060	0.060	0.058	0.060
$\beta_{112} = 1$	0.075	0.075	0.068	0.069	0.069	0.069	0.069	0.069	0.068	0.069
$\beta_{121} = 0.5$	0.096	0.094	0.075	0.077	0.075	0.077	0.077	0.078	0.075	0.077
$\beta_{122} = -2$	0.055	0.063	0.054	0.057	0.054	0.057	0.054	0.057	0.054	0.057
$\beta_{211} = 1.75$	0.264	0.374	0.087	0.121	0.058	0.088	0.487	0.135	0.045	0.045
$\beta_{212} = 0.25$	0.182	0.316	0.116	0.166	0.090	0.127	0.131	0.142	0.073	0.074
$\beta_{221} = 1$	0.211	0.339	0.096	0.120	0.080	0.105	0.111	0.117	0.067	0.073
$\beta_{222} = 1$	0.437	0.548	0.126	0.181	0.113	0.130	0.440	0.341	0.103	0.111
$\beta_{311} = 1.75 + \epsilon$	0.089	0.176	0.083	0.100	0.074	0.089	0.074	0.085	0.059	0.063
$\beta_{312} = 0.25 + \epsilon$	0.185	0.252	0.130	0.167	0.117	0.152	0.125	0.153	0.104	0.127
$\beta_{321} = 1 + \epsilon$	0.112	0.275	0.103	0.115	0.083	0.113	0.086	0.099	0.076	0.088
$\beta_{322} = 1 + \epsilon$	0.149	0.197	0.131	0.150	0.125	0.176	0.132	0.145	0.119	0.135
Irrelevant regres	sors									
$\beta_{113} = 0$	_	0.057	_	0.048	_	0.048	_	0.052	_	0.048
$\beta_{123} = 0$	_	0.082	_	0.081	_	0.081	_	0.080	_	0.080
$\beta_{213} = 0$	_	0.179	_	0.106	_	0.099	_	0.081	_	0.062
$\beta_{223} = 0$	_	0.214	_	0.200	_	0.156	_	0.126	_	0.125
$\beta_{313} = 0$	_	0.353	_	0.139	_	0.108	_	0.101	_	0.089
$\beta_{323} = 0$	_	0.294	_	0.163	_	0.172	_	0.162	_	0.163

Table 4: Summary statistics of the ARI index under SuCW and CW models in the first study: mean, median, interquartile range (IQR) and standard deviation (SD). The p-values in the last row refer to the paired samples Wilcoxon test for the hypothesis of equality between ARIs for each ϵ .

	$\epsilon = 0$	0.275	$\epsilon =$	0.3	$\epsilon = 0$	0.325	$\epsilon = 0$	0.350	$\epsilon = 0$	0.375
	SuCW	CW	SuCW	CW	SuCW	CW	SuCW	CW	SuCW	CW
Mean	0.852	0.806	0.901	0.887	0.927	0.920	0.936	0.936	0.956	0.956
Median	0.878	0.862	0.911	0.901	0.930	0.929	0.945	0.947	0.959	0.960
IQR	0.062	0.192	0.035	0.043	0.030	0.034	0.026	0.028	0.020	0.023
SD	0.083	0.121	0.037	0.070	0.026	0.048	0.057	0.053	0.018	0.019
p-value	< 1	0^{-5}	0.0	717	0.5	573	0.2	262	0.9	35

Table 5: Distributions of the number of components for the best SuCW and CW models in the second study.

	$\epsilon = 0.$	275	$\epsilon = 0.3$		$\epsilon = 0.$	$\epsilon = 0.325$		350	$\epsilon = 0.375$	
	SuCW	CW	SuCW	CW	SuCW	CW	SuCW	CW	SuCW	CW
K=2	48	99	7	84	1	39	1	10	0	1
K = 3	52	1	93	16	99	61	98	90	100	99
K = 4	0	0	0	0	0	0	1	0	0	0

the standard deviation of ARI seem to show a decreasing trend. Furthermore, SuCW models tend to be characterised by a greater ability to properly estimate 493 the true classification of the sample observations for each examined value of ϵ , 494 even though the differences in terms of mean and median values of ARI seem 495 to vanish for larger values of ϵ . This pattern is confirmed by the results of the 496 paired samples Wilcoxon test, suggesting that the difference between the mean values of ARI with the two approaches appears to be statistically significant 498 only when the degree of separation is low (see the p-values in the last row of 499 Table 4). This behaviour might be connected to the fact that, as the value of ϵ 500 increases, not only the differences between the two approaches in recovering the 501 actual values of the parameters tend to vanish, but also there is an improvement in the ability of CW models to recognise the presence of irrelevant regressors. 503

A second analysis has been carried out, where the performance of the two 504 approaches has been evaluated without exploiting the knowledge of neither the 505 number of components nor the parameterisation of the component-covariance matrices of X and Y employed to generate the datasets. Thus, 597 different 507 SuCW models have been estimated for each simulated dataset: 196 differentially 508 parameterised models for each K = 2, 3, 4 and 9 models with K = 1. The same 509 task has been carried out by employing CW models. Then, the best SuCW and CW 510 models fitted to each dataset have been selected according to the BIC. Table 5 511 summarises the results of this procedure in terms of recovery of the true K. 512 The impact of the value of ϵ on this aspect is evident. Generally speaking, the 513 ability to select the correct value of K improves as the separation increases. 514 By focusing the attention on the distributions of the number of components 515 for the best CW models fitted to the 100 datasets, it emerges that with such an approach the true number of components tends to be severely underestimated 517 with the two lowest levels of separation ($\epsilon = 0.275, 3$). On the contrary, using 518 SuCW models leads to the selection of the correct number of components for the 519 majority of the simulated datasets with all levels of separation; furthermore, the 520 proportion of datasets for which the selected SuCW model has three components 521 increases quickly with ϵ , reaching 93% when $\epsilon = 0.3$ and approaching nearly 522 100% for larger values of ϵ . 523

In order to assess the possible consequences of a wrong choice of K on 524 the ability of CW models to recognise the presence of equation-specific irrelevant regressors, the biases of the estimates of the effects of these regressors have been computed for CW models with K equal to 1, 2 and 4. According to the values 527 reported in Table 6, it appears that the estimates of the regression coefficients for 528 the irrelevant regressors can be severely biased when the number of components 529 is lower than the true one. On the contrary, when the number of components 530 exceeds the true K, the results are comparable with those obtained in the first analysis using models with K=3 components (see the lower part of Table 2). 532 It is also worth noting that for some coefficients the bias seems to show a trend 533 which increases with the separation among components.

Table 6: Bias for the regression coefficients for equation-specific irrelevant regressors under ${\tt CW}$ models in the second study.

		$\epsilon = 0.275$	$\epsilon = 0.3$	$\epsilon = 0.325$	$\epsilon = 0.350$	$\epsilon = 0.375$
K = 1	$\beta_{113} = 0$	0.047	0.065	0.084	0.105	0.127
K = 1	$\beta_{123} = 0$	0.525	0.504	0.483	0.462	0.440
	$\beta_{113} = 0$	0.003	0.003	0.003	0.003	0.003
K = 2	$\beta_{123} = 0$	0.022	0.022	0.022	0.022	0.022
K = Z	$\beta_{213} = 0$	0.377	0.428	0.479	0.529	0.578
	$\beta_{223} = 0$	0.110	0.128	0.147	0.166	0.186
	$\beta_{113} = 0$	0.011	0.024	0.023	0.008	0.032
	$\beta_{123} = 0$	0.034	0.047	0.043	0.035	0.027
	$\beta_{213} = 0$	0.017	0.012	0.004	0.014	0.013
K = 4	$\beta_{223} = 0$	0.095	0.015	0.011	0.093	0.065
N=4	$\beta_{313} = 0$	0.056	0.017	0.010	0.011	0.012
	$\beta_{323} = 0$	0.009	0.056	0.004	0.011	0.004
	$\beta_{413} = 0$	0.070	0.017	0.026	0.024	0.018
	$\beta_{423} = 0$	0.057	0.024	0.013	0.013	0.000

As far as the classification recovery is concerned, the obtained results demon-535 strates that the ability to estimate the true classification of the sample observa-536 tions with both approaches increases with ϵ . However, the gap between the two approaches in terms of mean and median ARI is quite large and statistically 538 significant for the three smallest values of ϵ (see the Table 7). It is worth noting 539 that the behaviour of the variability of the ARI index is strictly related to the 540 variability in the distribution of the optimal value of K selected according to the BIC. In summary, the obtained results seem to suggest that the inclusion of the regressor X_3 in the equation (15) and the regressor X_2 in the equation (16) has 543 a negative impact both on the choice of the correct number of components and 544 on the reconstruction of the true classification of the sample observations. How-545 ever, the consequences of including these irrelevant regressors seem to become negligible as the separation among components increases. A possible explanation of this behaviour could be related to the fact that the clustering task is 548 eased when the components are well-separated. In such situations, even if CW 549 models are non-parsimonious, they can lead to the correct choice of K. As a 550 consequence, they are able to provide estimates for the regression coefficients of 551 irrelevant regressors that are sufficiently close to zero, so that the inclusion of 552 such regressors has little effect on the estimated posterior probabilities employed 553 to classify the sample observations. 554

55 4. Results from the analysis of real data

Two real situations have been examined to evaluate the practical usefulness of SuCW models in comparison with CW models. For both these model classes, models have been estimated for K from 1 to 9. For each of these values, all possible parsimonious CW and SuCW models have been fitted (see Section 2.7). Analyses of the examined real datasets have been carried out also through the clusterwise regression models described in Section 2.2. Namely, the comparison with models (5) allows to assess the adequacy of the assignment independence assumption. Furthermore, from the comparison with models (6) it is possi-

Table 7: Summary statistics of the ARI index for the best model SuCW and CW model in the second study: mean, median, interquartile range (IQR) and standard deviation (SD). The p-values in the last row refer to the paired samples Wilcoxon test for the hypothesis of equality between ARIs for each ϵ .

	$\epsilon = 0$	0.275	$\epsilon = 0.3$		$\epsilon = 0$	0.325	$\epsilon = 0$	0.350	$\epsilon = 0$	0.375
	SuCW	CW	SuCW	CW	SuCW	CW	SuCW	CW	SuCW	CW
Mean	0.766	0.644	0.885	0.687	0.924	0.820	0.939	0.916	0.956	0.953
Median	0.785	0.644	0.905	0.652	0.929	0.904	0.945	0.945	0.959	0.960
IQR	0.236	0.036	0.041	0.048	0.030	0.273	0.026	0.030	0.020	0.023
SD	0.120	0.034	0.071	0.103	0.035	0.138	0.038	0.087	0.018	0.036
p-value	< 10	0^{-10}	< 10	0^{-15}	< 1	0^{-8}	0.0)22	0.8	351

ble to establish whether fitting multivariate cluster-weighted models based on 564 seemingly unrelated linear regression for D responses leads to an improvement 565 over an approach based on D univariate seemingly unrelated linear clusterwise 566 regression models. From now on, models (5) and (6) are denoted as SuCR and uSuCR, respectively. Models from equations (5) and (6) have been estimated 568 also using the same vector of covariates for all responses (i.e., with $\mathbf{x}_{id} = \mathbf{x}_i$ 569 $\forall d$); in the following, they are denoted as CR and uCR, respectively. All these 570 clusterwise regression models have been fitted for a number of components from 571 1 to 9 through a specific function developed in the R environment which also 572 allows the estimation of seemingly unrelated linear parsimonious clusterwise 573 models (for more details see Galimberti and Soffritti, 2020). Parameters $\mu_{\mathbf{X}}$ 574 and $\Sigma_{\mathbf{X}}$ of the Gaussian distribution for the covariates in models (5) and (6) 575 have been estimated under three possible structures of $\Sigma_{\mathbf{X}}$: fully unconstrained 576 (VV), diagonal with P unequal variances (VI) and diagonal with equal variances 577 (EI). As far as the variances $\sigma_{k_d d}$, $k_d = 1, \dots, K_d$, in the univariate clusterwise 578 regression models are concerned, the estimation has been carried out under both 579 an homoscedastic (E) and heteroscedastic (V) assumption.

4.1. Canned tuna sales in USA

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Data taken from Chevalier et al. (2003) and available within the R package 582 bayesm (Rossi, 2012) provides information about seven of the top 10 U.S. brands in the canned tuna product category for I = 338 weeks between September 584 1989 and May 1997 (tuna dataset). The available information is the volume 585 of weekly sales (Move), a measure of the display activity (Nsale) and the log 586 price (Lprice) of each brand. Analyses illustrated here have been focused on 587 D=2 products: Bumble Bee Chunk 6.12 oz. (BBC) and Bumble Bee Solid 6.12 oz. (BBS). A previous study about the effect of prices and promotional 589 activities on sales for these two products, based on clusterwise linear regression 590 models (Galimberti and Soffritti, 2020), demonstrated that the effect of log price 591 on log unit sales is not homogeneous during the examined period of time for 592 both products. Furthermore, a search for the predictors to be employed in the two regression equations showed that models including only the log unit prices 594 should be preferred. Thus, the analysis here has been focused on four variables: 595 $\mathbf{Y} = (\texttt{Lmove BBC}, \texttt{Lmove BBS})', \mathbf{X} = (\texttt{Lprice BBC}, \texttt{Lprice BBS})', \text{ where Lmove}$ 596 denotes the logarithm of Move. As typically happens with food prices, also 597 prices of BBC and BBS appear to change according to an almost discrete grid of 598 values (see the scatterplot on the left part of Figure 3). Although the Pearson's 599 correlation coefficient between the two responses is low (0.1844), according to 600 the Student's t test the hypothesis of linear independence between Lmove BBC 601 and Lmove BBS has to be rejected; Lmove BBC results to be negatively and strongly correlated with Lprice BBC; there is also a negative and significant linear dependence between Lmove BBS and the logarithm of the prices for both 604 products (see Table 8). 605

By assuming that prices for each of the two examined products can only 606 affect sales of the same product, SuCW, SuCR and uSuCR models have been specified by using Lprice BBS as regressor in the equation for Lmove BBS, Lprice BBC as regressor for Lmove BBC. Table 9 reports the models which best fit the 609 tuna dataset according to the BIC for each combination of the nine examined 610 values of K and each of the model types SuCW and CW. All these models have

Table 8: Pearson's correlation matrix (lower diagonal part) and p-values of the Student's t test for the hypothesis of linear independence between variables (upper diagonal part) from the tuna dataset.

	Lmove BBC	Lmove BBS	Lprice BBC	Lprice BBS
Lmove BBC	1.0000	0.0007	$< 10^{-67}$	0.2678
Lmove BBS	0.1844	1.0000	0.0011	$< 10^{-8}$
Lprice BBC	-0.7727	-0.1767	1.0000	0.4420
Lprice BBS	-0.0604	-0.3172	0.0420	1.0000

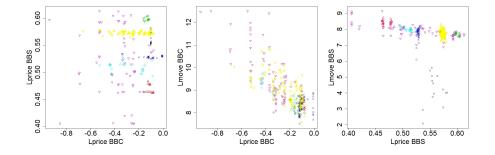


Figure 3: Scatterplots for three pairs of variables from the analysis of the tuna dataset. Observations are pictured with seven different colours and symbols according to the classification obtained from the best model.

Table 9: Best models fitted to the tuna dataset within some given model classes and their BIC values. Columns acr.X and acr.Y report the acronyms of the parsimonious paramaterisations for the component-covariance matrices of \mathbf{X} and \mathbf{Y} , respectively.

Model	K	acr.X	acr.Y	BIC_M	Model	K	acr.X	acr.Y	BIC_M
SuCW	1	EEI	EEI	-18.9	CW	1	EEI	EEI	-18.1
SuCW	2	VVE	EVV	-812.2	CW	2	VVE	EVV	-794.2
SuCW	3	VVI	EVV	-929.1	CW	3	VEV	VVE	-922.3
SuCW	4	VVE	VVE	-1195.0	CW	4	VVE	VVE	-1157.4
SuCW	5	VVI	VEV	-1282.0	CW	5	VVI	VVE	-1267.1
SuCW	6	VVI	VEV	-1355.2	CW	6	VVI	VVE	-1333.7
SuCW	7	VVI	VEV	-1389.8	CW	7	VVI	VVE	-1331.4
SuCW	8	VVI	VEV	-1387.2	CW	8	VVI	VVE	-1341.3
SuCW	9	VVI	VEV	-1371.1	CW	9	VVI	VVI	-1326.4

been estimated within a limit of 237 iterations of the ECM algorithm. Figure 4 shows the values of the BIC for the best CW and SuCW models by K. As far 613 as the clusterwise regression models are concerned, Table 10 summarises some 614 information about the best fitted models within each of the model classes SuCR, 615 CR, uSuCR and uCR obtained from equations (5) and (6). Overall, it seems that the best trade-off between the fit and complexity can be obtained using the SuCW 617 model with K = 7 clusters of weeks. The convergence of the ECM algorithm for 618 the parameter estimation has been reached after 53 iterations. For the clusters 619 detected by this model, the distributions of the two regressors are diagonal with variable volumes and shapes. As far as the joint conditional distributions of 62 the two responses given the corresponding regressors are concerned, clusters are 622 characterized by ellipsoidal distributions with variable volumes and orientations 623 and equal shape. 624

The first cluster is composed of 16 consecutive weeks corresponding to the period from end-October 1990 to mid-February 1991 (see the additional information about this dataset available at the University of Chicago website http://research.chicagobooth.edu/kilts/marketing-databases/dominicks). In that

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Table 10: Best models fitted to the tuna dataset within the model classes defined from equations (5) and (6) and their BIC values.

Model	Best fitted model	BIC_M
$\phi_{P}\left(\mathbf{x}_{i};oldsymbol{\mu}_{\mathbf{X}},oldsymbol{\Sigma}_{\mathbf{X}} ight)$	M_a : $acr.X = VI$	-1408.0
$\sum_{k=1}^{K} \pi_k \phi_D\left(\mathbf{y}_i \mathbf{x}_i; \mathcal{X}_i' \boldsymbol{\beta}_k^*, \boldsymbol{\Sigma}_{\mathbf{Y}_k}\right)$	M_b : $K=3$, $acr.Y={\tt EVV}$	652.5
$\sum_{k=1}^{K} \pi_k \phi_D\left(\mathbf{y}_i \mathbf{x}_i; \mathbf{B}_k'\mathbf{x}_i^*, \mathbf{\Sigma}_{\mathbf{Y}_k}\right)$	M_c : $K = 3$, $acr.Y = VVE$	683.4
$\sum_{k=1}^{K} \pi_k \phi_1 \left(y_{i1} \mathbf{x}_i; \mathbf{x}_{i1}^{*'} \boldsymbol{\beta}_{k1}^*, \sigma_{k1}^2 \right)$	M_d : $K = 2$, $acr.Y = V$	496.1
$\sum_{k=1}^{K} \pi_k \phi_1 \left(y_{i1} \mathbf{x}_i; \boldsymbol{\beta}_{k1}^{*\prime} \mathbf{x}_i^*, \sigma_{k1}^2 \right)$	M_e : $K = 2$, $acr.Y = V$	505.4
$\sum_{k=1}^{K} \pi_k \phi_1 \left(y_{i2} \mathbf{x}_i; \mathbf{x}_{i2}^{*'} \boldsymbol{\beta}_{k2}^*, \sigma_{k2}^2 \right)$	M_f : $K=2$, $acr.Y=V$	162.0
$\sum_{k=1}^{K} \pi_k \phi_1 \left(y_{i2} \mathbf{x}_i; \boldsymbol{\beta}_{k2}^{*\prime} \mathbf{x}_i^*, \sigma_{k2}^2 \right)$	M_g : $K = 2$, $acr.Y = V$	164.7
SuCR	M_a and M_b	-755.5
CR	M_a and M_c	-724.6
uSuCR	M_a , M_d and M_f	-749.9
uCR	M_a , M_e and M_g	-737.9

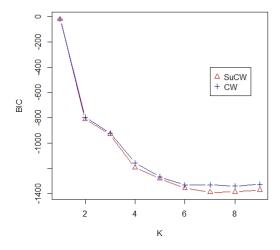


Figure 4: Tuna dataset: BIC values of the best ${\tt CW}$ and ${\tt SuCW}$ models by number of components.

Table 11: Estimated π_k , $\mu_{\mathbf{X}_k}$ and $\boldsymbol{\beta}_k^*$ of the best model fitted to the tuna dataset. $\mathbf{a}[l]$ denotes the lth element of vector \mathbf{a} .

k	1	2	3	4	5	6	7
$\hat{\pi}_k$	0.047	0.063	0.080	0.098	0.104	0.254	0.354
$\hat{m{\mu}}_{\mathbf{X}_k}[1]$	-0.103	-0.106	-0.130	-0.085	-0.297	-0.366	-0.245
$\hat{\boldsymbol{\mu}}_{\mathbf{X}_k}[2]$	0.554	0.468	0.599	0.528	0.511	0.520	0.573
$\hat{\boldsymbol{\beta}}_{k1}^*[1]$	7.946	8.506	8.925	8.203	7.844	7.751	8.239
$\hat{\boldsymbol{\beta}}_{k1}^*[2]$	-4.493	0.801	5.187	-0.421	-3.544	-5.090	-3.173
$\hat{\boldsymbol{\beta}}_{k2}^*[1]$	17.051	8.820	16.503	-4.040	13.142	10.504	20.766
$\hat{\boldsymbol{\beta}}_{k2}^*[2]$	-22.692	-0.855	-14.724	22.436	-10.115	-4.920	-22.712

period a worldwide boycott campaign (promoted by the U.S. nongovernmental 629 organisation Earth Island Institute) encouraged consumers not to buy Bumble 630 Bee tuna because Bumble Bee was found to be buying yellow-fin tuna caught by dolphin-unsafe techniques (Baird and Quastel, 2011). The negative impact 632 of such a campaign on Bumble Bee tuna sales appears to be evident for BBS 633 (see the black points in the scatterplot of Figure 3 for this product). The mean 634 prices of both products in the weeks of this cluster are quite high (see the first column in Table 11). Furthermore, prices of BBC in this cluster are highly 636 homogeneous, as suggested by the low variance of Lprice BBC (not reported 637 here). Finally, the effect of prices on sales in the same weeks is negative and 638 particularly strong for BBS (see Table 11). The second cluster comprises 22 639 weeks (red points in the scatterplots of Figure 3), some of which are in close correspondence with Easter 1990 and 1991, Christmas 1993, Presidents day and Labor day 1994. They are mainly characterized by the lowest mean price of BBS 642 and a negligible impact of prices on sales for both products. Furthermore, prices 643 of BBS in such weeks result to be quite homogeneous. Cluster 3 is composed 644 of 27 weeks (green points in Figure 3) with the highest mean price of BBS. In this cluster, the effects of prices on sales are negative for BBS and positive for BBC; furthermore, prices of both products are homogeneous. The special events 647 corresponding to the weeks of cluster 3 are: Memorial days 1994 and 1995, 4th 648

of July 1994 and 1995, Halloween and Thanksgiving 1994. Cluster 4 is mainly composed of weeks from end-November 1995 to end-April 1997; two distinctive 650 features of this cluster (34 weeks, dark blue points) are that it shows the highest mean price of BBC and highly homogeneous prices of BBS; furthermore, the 652 estimated effect of prices on sales of BBS is positive and particularly strong. 653 Labor day 1991, January 1992, Memorial days 1992 and 1993 are the events 654 and periods associated with the weeks in cluster 5 (37 weeks, sky-blue points), 655 which is characterized by intermediate mean prices and mild negative effects of prices on sales for both products. As far as clusters 6 and 7 are concerned, 657 they contain 78 (purple points) and 124 (yellow points) weeks, respectively. The 658 main distinctive feature of cluster 6 is that the variances of Lprice BBC and 659 Lprice BBS (not reported here) are extremely large; furthermore, this cluster registers the lowest mean price of BBC. Cluster 6 mainly comprises weeks from mid-September 1991 to end-December 1991, January 1993, and the periods as-662 sociated with Christmas 1992, Presidents day 1992 and 1993, Easter 1992, 1993 663 and 1995. Weeks belonging to cluster 7 are characterized by high and highly 664 homogeneous prices of BBS; furthermore, the effect of prices on sales of BBS in 665 these weeks is negative and particularly strong. In summary, by focusing the attention on the estimated regression coefficients of the seven clusters of weeks 667 detected by the model, the main interesting findings are a clear evidence of dif-668 ferential effects of the log prices on the log unit sales for both products and the 669 identification of two clusters in which such effects are positive for either BBS or BBC. The overall agreement between this partition and the one produced 67 by the best CW model, which is composed of 8 clusters (see Table 12), is high 672 (ARI = 0.8293): weeks have been classified in almost the same way by the two 673 approaches; some exceptions mainly involve the sixth cluster of the partition 674 illustrated above. 675

The comparison between these results and those produced from the best fitted linear clusterwise regression model (see the SuCR model in Table 10) shows that in the analysed dataset there is an additional source of heterogeneity over time, which appears to lie mainly in the prices of BBC tuna. Thus, when

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Table 12: Cross-classification of the observations from the tuna dataset, based on the maximum posterior probabilities estimated from the best CW and SuCW fitted models. Labels for clusters reported in rows and colums refer to CW and SuCW, respectively.

k	1	2	3	4	5	6	7
1	0	0	0	0	0	6	0
2	16	0	0	0	0	0	0
3	0	21	0	0	0	0	0
4	0	0	27	0	0	0	0
5	0	0	0	33	0	1	0
6	0	1	0	1	36	21	0
7	0	0	0	0	1	48	2
8	0	0	0	0	0	2	122

modelling the joint distribution of prices and sales for both products, more 680 clusters have been detected (7 instead of 3). A further difference between the results obtained from these two approaches is that all the effects of log prices on the log unit sales for both products results to be negative within each cluster 683 identified by the best linear clusterwise regression model. It is also worth noting 684 that there is an almost perfect correspondence between one of the three clusters 685 identified through the best linear clusterwise regression model and the first 686 cluster described above (see Galimberti and Soffritti, 2020, for more details on the results obtained from the analysis of these data through the clusterwise 688 regression approach). 689

As illustrated in Section 2.1, an underlying assumption of the best fitted 690 model is that both $\mathbf{X}|\Omega_k$ and $\mathbf{Y}|(\mathbf{X}=\mathbf{x},\Omega_k)$ follow a multivariate normal distribution for k = 1, ..., K. An evaluation of the adequacy of such an assumption for the examined dataset has been carried out by resorting to some measures 693 of multivariate skewness and kurtosis (Mardia, 1970, 1974); by exploiting their asymptotic distribution derived under the hypothesis of multivariate nor-695 mality, those measures can also be employed as statistics for testing the hypothesis of multivariate normality. Namely, the function mult.norm of the R

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Table 13: P-values of Mardia's skewness and kurtosis statistics for the residuals $\mathbf{x} - \boldsymbol{\mu}_{\mathbf{X}_k}$ and $\mathbf{y} - \mathcal{X}'\boldsymbol{\beta}_k, \, k = 1, \dots, K$, computed from the best model fitted to the tuna dataset.

k	1	2	3	4	5	6	7
$\mathbf{x} - \boldsymbol{\mu}_{\mathbf{X}_k}$							
skewness	0.2654	0.1887	0.0124	0.0370	0.7230	0.0013	0.0000
kurtosis	0.9963	0.3052	0.6470	0.3134	0.0510	0.0777	0.0371
$\mathbf{y} - \mathcal{X}' oldsymbol{eta}_k$							
skewness	0.0086	0.3526	0.7434	0.0829	0.8385	0.0002	0.0185
kurtosis	0.1308	0.5668	0.2121	0.3235	0.0483	0.0008	0.3927

package QuantPsych (Fletcher, 2012) has been employed to compute the values 698 of such measures within each cluster detected by the best model from the es-699 timated residuals $\mathbf{x}_i - \hat{\boldsymbol{\mu}}_{\mathbf{X}_k}$ and $\mathbf{y}_i - \mathcal{X}_i' \hat{\boldsymbol{\beta}}_k \ \forall (i,k) \in \{(i,k), i \in \{1,\dots,I\}, k = 1,\dots, k \}$ $\arg \max_{h} \{\hat{\tau}_{ih}, h = 1, \dots, K\}\};$ the p-values associated with the so obtained results are summarised in Table 13. Based on these findings, in the first five 702 clusters the null hypothesis of multivariate normality should not be rejected 703 at a Bonferroni-corrected 0.05/7 = 0.0071 significance level. On the contrary, 704 both types of residuals clearly deviate from the multivariate normality within 705 the sixth cluster. As far as the seventh cluster is concerned, the null hypothesis should be rejected only for the residuals $\mathbf{x} - \boldsymbol{\mu}_{\mathbf{X}_k}$. 707

4.2. Regional tourism in Italy

In line with studies aiming at evaluating the link between tourism flows and attendance at museums and monuments (see, e.g., Cellini and Cuccia, 2013), the data analysed here provides information about tourist arrivals (denoted Arriv), tourist overnights (Overn) and visits to State museums, monuments and museum networks (Visit) with a monthly frequency over the period January 1999 to December 2017 in two Italian regions: Emilia Romagna (ER) and Veneto (Ve).

Data concerning Visit has been obtained from the website of the Italian Min-

istry of Cultural Heritage¹; the sources for Arriv and Overn are the websites of the two regional governments². In this dataset the average stays (AvStay), 717 computed as the ratio between Overn and Arriv, are also provided. Thus, the 718 dataset is composed of I = 228 monthly observations for eight variables; from 719 now on, it has been denoted as RtI. The goal of the analysis is to study the effect 720 of the tourist arrivals and average stays on the visits to State museums, mon-721 uments and museum networks in Emilia-Romagna and Veneto. Thus, in this 722 analysis Y = (Visit ER, Visit Ve)', X = (Arriv ER, AvStay ER, Arriv Ve,AvStay Ve)'. The analysis has been performed using data in thousands. Fig-724 ure 5 shows the bivariate scatterplots for pairs of regressors and pairs composed 725 of one response and one regressor; month abbreviations are used as labels for the 726 observations. Visits to to State museums, monuments and museum networks in 727 the two regions result to be highly linearly dependent (see Table 14); high and positive pairwise correlations also characterise tourist arrivals and average stays 729 in either region; the hypothesis of linear independence is not rejected between 730 Visit ER and the average stays; the same result holds true also for Visit Ve. 731

Table 14: Pearson's correlation matrix (lower diagonal part) and p-values of the Student's t test for the hypothesis of linear independence between variables (upper diagonal part) from the RtI dataset.

	Visit ER	Visit Ve	Arriv ER	AvStay ER	Arriv Ve	AvStay Ve
Visit ER	1.0000	$< 10^{-66}$	0.0002	0.3684	0.0003	0.3879
Visit VE	0.8562	1.0000	$< 10^{-8}$	0.7807	$< 10^{-9}$	0.9572
Arriv ER	0.2421	0.3722	1.0000	$< 10^{-53}$	$< 10^{-166}$	$< 10^{-47}$
AvStay ER	-0.0598	0.0185	0.8081	1.0000	$< 10^{-45}$	$< 10^{-175}$
Arriv Ve	0.2394	0.4015	0.9826	0.7696	1.0000	$< 10^{-40}$
AvStay Ve	-0.0575	-0.0036	0.7833	0.9856	0.7456	1.0000

http://www.statistica.beniculturali.it.

https://www.veneto.eu/web/area-operatori/statistiche.

²https://statistica.regione.emilia-romagna.it/turismo,

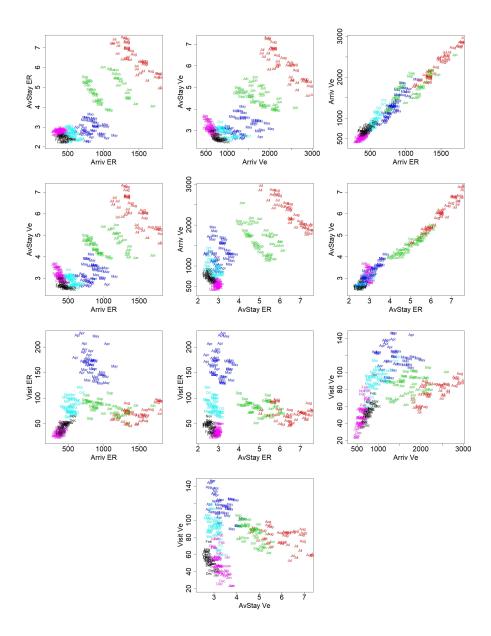


Figure 5: Bivariate scatterplots for pairs of variables in the analysis of the RtI dataset. Month abbreviations are used as labels. Observations are coloured according to the classification obtained from the best model.

Table 15: Best models fitted to the RtI dataset within some given model classes and their BIC values. Columns acr.X and acr.Y report the acronyms of the parsimonious paramaterisations for the component-covariance matrices of \mathbf{X} and \mathbf{Y} , respectively.

Model	K	acr.X	acr.Y	BIC_M	Model	K	acr.X	acr.Y	BIC_M
SuCW	1	EEE	EEE	10948.6	CW	1	EEE	EEE	10965.7
SuCW	2	VVV	EEV	10223.8	CW	2	VVV	EEV	10146.2
SuCW	3	VVV	VVV	9991.5	CW	3	VVV	VVE	9934.6
SuCW	4	VVV	VEV	9898.8	CW	4	VVV	VEE	9886.6
SuCW	5	EVV	VEE	9822.3	CW	5	VVV	VII	9751.2
SuCW	6	VVV	VVI	9716.5	CW	6	VVV	VEE	9788.3
SuCW	7	VVV	VEV	9736.9	CW	7	VVV	VEE	9799.4
SuCW	8	EVV	VEV	9796.0	CW	8	EVV	VEE	9861.2
SuCW	9	VVV	VEV	9815.0	CW	9	VVV	VEE	9917.1

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A first analysis has been performed by assuming that arrivals and average stays in each of the two regions can only affect attendance at museums and monuments of the same region. Thus, SuCW models have been specified by using Arriv ER and AvStay ER as regressors in the equation for Visit ER, Arriv Ve and AvStay Ve as regressors for Visit Ve. However, since Emilia-Romagna and Veneto are neighboring regions, arrivals and average stays in one region could also have an impact on the visits to State museums and monuments of the other region, hence the second analysis has been carried out through CW models. Table 15 provides information about the models which best fit the RtI dataset according to the BIC for each combination of the nine examined values of K and the two fitted model types. The convergence of the ECM algorithm for the estimation of these models has been reached within a limit of 161 iterations. Figure 6 shows the values of the BIC for the best CW and SuCW models by K. Table 16 provides a summary of the results obtained from the best fitted models within each of the model classes defined by equations (5) and (6). Overall, the model with the best trade-off between the fit and complexity seems to be the SuCW model with K=6 clusters of months. The ECM algorithm for the

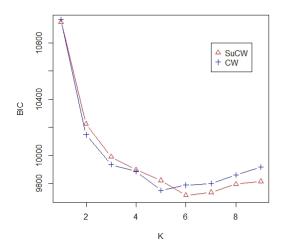


Figure 6: RtI dataset: BIC values of the best CW and SuCW models by number of components.

resulting clusters, the four regressors have ellipsoidal distributions with variable volumes, shapes and orientations. As far as the joint conditional distributions of the two responses given the corresponding regressors are concerned, clusters show diagonal distributions with variable volumes and shapes, suggesting that Visit Ve and Visit ER are independent, conditionally on the regressors and cluster membership.

Four clusters are perfectly related to some months (see Table 17). They are:

757 cluster 2: observations in July and August;

cluster 3: observations in June and September;

759 cluster 4: observations in April and May;

cluster 5: observations in March and October.

As far as the months from November to February are concerned, observations

from January 1999 to November 2010 and those of January 2011 and February

₇₆₃ 2012 have been assigned to cluster 6; cluster 1 comprises all the remaining

observations in such months. The obtained cluster structure clearly reflects

seasonal patterns characterising tourism flows. Observations in cluster 2 (July 765 and August) are characterized by the highest mean values of tourist arrivals 766 and average stays in both regions, followed by those in cluster 3 (June and September) and cluster 4 (April and May) (see Table 18). From the comparison 768 between clusters 1 and 6 it emerges that wintertime tourism flows have changed 769 in both regions, showing an increase in the mean number of arrivals and a 770 decrease in the mean number of stays in recent years (cluster 1). In all clusters, 771 Veneto is characterised by mean values of both regressors which are higher than 772 those of Emilia-Romagna except for the average stays from June to September. 773 As far as the estimated regression coefficients are concerned (see Table 18), the 774 first interesting finding is that the effects of both the tourist arrivals and the 775 average stays on the number of visits result to be not homogeneous during the examined period of time. In both regions, such effects are positive in July and August: in Emilia-Romagna, this result also holds true in the months belonging 778 to cluster 6. In the other clusters of months the effect of tourist arrivals are 779 generally positive in both regions, while the average number of stays seem to 780 have a negative impact on the number of visits. This latter impact in Veneto 781 appears to be stronger than that in Emilia-Romagna in April, May, June and 782 September; the opposite result holds true for all the other months. 783

The comparison between this partition and the one based on the maximum 784 posterior probabilities estimated from the best CW fitted model (see Table 19) 785 suggests that they are quite similar (ARI = 0.8014); the main difference is that according to the approach based on CW models all the observations in the 787 months from November to February should be grouped into the same cluster. 788 This latter result mainly depends on the fact that, in the best model fitted to the 789 RtI dataset within the class of CW models with K=6, the effects of both Arriv 790 Ve on Visit ER and Arriv ER on Visit Ve in two clusters have been estimated to be quite similar (detailed results are not reported) and, thus, a better tradeoff between the fit and the complexity is reached by the best fitted CW model with 793 K=5. Furthermore, the comparison between the results obtained through SuCW 794 models and those produced from linear clusterwise regression analyses (see Table

Table 16: Best models fitted to the RtI dataset within the model classes defined from equations (5) and (6) and their BIC values.

Model	Best fitted model	BIC_M
$\phi_{P}\left(\mathbf{x}_{i};oldsymbol{\mu}_{\mathbf{X}},oldsymbol{\Sigma}_{\mathbf{X}} ight)$	M_a : $acr.X = VV$	6808.1
$\sum_{k=1}^{K} \pi_k \phi_D\left(\mathbf{y}_i \mathbf{x}_i; \mathcal{X}_i'\boldsymbol{\beta}_k^*, \mathbf{\Sigma}_{\mathbf{Y}_k}\right)$	M_b : $K = 4$, $acr.Y = VEV$	3943.3
$\sum_{k=1}^{K} \pi_k \phi_D\left(\mathbf{y}_i \mathbf{x}_i; \mathbf{B}_k'\mathbf{x}_i^*, \mathbf{\Sigma}_{\mathbf{Y}_k}\right)$	M_c : $K=3$, $acr.Y=$ VEV	3920.2
$\sum_{k=1}^{K} \pi_k \phi_1 \left(y_{i1} \mathbf{x}_i; \mathbf{x}_{i1}^{*'} \boldsymbol{\beta}_{k1}^*, \sigma_{k1}^2 \right)$	M_d : $K = 4$, $acr.Y = V$	2166.6
$\sum_{k=1}^{K} \pi_k \phi_1 \left(y_{i1} \mathbf{x}_i; \boldsymbol{\beta}_{k1}^{*\prime} \mathbf{x}_i^*, \sigma_{k1}^2 \right)$	M_e : $K = 3$, $acr.Y = V$	2183.4
$\sum_{k=1}^{K} \pi_{k} \phi_{1} \left(y_{i2} \mathbf{x}_{i}; \mathbf{x}_{i2}^{*'} \boldsymbol{\beta}_{k2}^{*}, \sigma_{k2}^{2} \right)$	M_f : $K = 4$, $acr.Y = E$	1987.2
$\sum_{k=1}^{K} \pi_k \phi_1 \left(y_{i2} \mathbf{x}_i; \boldsymbol{\beta}_{k2}^{*\prime} \mathbf{x}_i^*, \sigma_{k2}^2 \right)$	M_g : $K = 2$, $acr.Y = V$	1987.1
SuCR	M_a and M_b	10751.4
CR	M_a and M_c	10728.3
uSuCR	M_a , M_d and M_f	10962.0
uCR	M_a , M_e and M_g	10978.6

Table 17: Cross-classification of the observations from the RtI dataset, based on their variable time identified by month and maximum posterior probability estimated from the best fitted model.

k	Jan	Feb	Mar	Apr	May	Jun	Jul	Aug	Sep	Oct	Nov	Dec
1	6	6	0	0	0	0	0	0	0	0	7	8
2	0	0	0	0	0	0	19	19	0	0	0	0
3	0	0	0	0	0	19	0	0	19	0	0	0
4	0	0	0	19	19	0	0	0	0	0	0	0
5	0	0	19	0	0	0	0	0	0	19	0	0
6	13	13	0	0	0	0	0	0	0	0	12	11

Table 18: Estimated π_k , $\mu_{\mathbf{X}_k}$ and $\boldsymbol{\beta}_k^*$ of the best model fitted to the RtI dataset. $\mathbf{a}[l]$ denotes the lth element of vector \mathbf{a} .

k	1	2	3	4	5	6
$\hat{\pi}_k$	0.121	0.167	0.167	0.167	0.167	0.213
$\hat{m{\mu}}_{\mathbf{X}_k}[1]$	377.1	1389.8	1001.0	767.7	487.4	316.5
$\hat{\boldsymbol{\mu}}_{\mathbf{X}_k}[2]$	2.350	6.304	4.847	2.823	2.572	2.694
$\hat{\boldsymbol{\mu}}_{\mathbf{X}_k}[3]$	665.5	2248.1	1639.0	1251.5	867.7	502.5
$\hat{\boldsymbol{\mu}}_{\mathbf{X}_k}[4]$	2.588	5.894	4.511	3.119	2.836	3.039
$\hat{oldsymbol{eta}}_{k1}^*[1]$	34.063	-263.846	143.648	278.219	163.170	-68.964
$\hat{\boldsymbol{\beta}}_{k1}^*[2]$	0.117	0.116	-0.028	-0.095	0.008	0.090
$\hat{\boldsymbol{\beta}}_{k1}^*[3]$	-15.948	26.302	-7.994	-14.227	-34.080	25.994
$\hat{\boldsymbol{\beta}}_{k2}^*[1]$	29.502	-43.971	183.051	179.415	106.489	-6.196
$\hat{\boldsymbol{\beta}}_{k2}^*[2]$	0.071	0.035	-0.006	0.003	0.032	0.132
$\hat{\boldsymbol{\beta}}_{k2}^*[3]$	-9.889	6.876	-20.067	-21.422	-16.044	-4.297

16) demonstrates that there is some clear evidence of seasonal heterogeneity not only in attendance at museums and monuments but also in tourism flows. Finally, a joint analysis for the two examined Italian regions based on seemingly unrelated cluster-weighted models results to be more effective than two separate linear clusterwise regression analyses.

As in the previous application, the mult.norm function of the R package QuantPsych has been employed to obtain an evaluation of the adequacy of the normality assumption within each cluster detected by the best model, based on the p-values of Mardia's measures of multivariate skewness and kurtosis computed from the estimated residuals $\mathbf{x}_i - \hat{\boldsymbol{\mu}}_{\mathbf{X}_k}$ and $\mathbf{y}_i - \mathcal{X}_i' \hat{\boldsymbol{\beta}}_k \ \forall (i,k) \in \{(i,k), i \in \{1,\ldots,I\}, k = \arg\max_h\{\hat{\tau}_{ih}, h = 1,\ldots,K\}\}$ (see Table 20. The obtained results suggest that the null hypothesis of multivariate normality should not be rejected at a Bonferroni-corrected 0.05/6 = 0.0083 significance level in any cluster

Table 19: Cross-classification of the observations from the RtI dataset, based on the maximum posterior probabilities estimated from the best CW and SuCW fitted models. Labels for clusters reported in rows and column refer to CW and SuCW, respectively.

k	1	2	3	4	5	6
1	0	0	34	0	0	0
2	0	0	0	38	0	0
3	0	0	0	0	38	0
4	0	38	4	0	0	0
5	27	0	0	0	0	49

Table 20: P-values of Mardia's skewness and kurtosis statistics for the residuals $\mathbf{x} - \boldsymbol{\mu}_{\mathbf{X}_k}$ and $\mathbf{y} - \mathcal{X}'\boldsymbol{\beta}_k, \, k = 1, \dots, K$, computed from the best model fitted to the RtI dataset.

k	1	2	3	4	5	6
$\mathbf{x} - \boldsymbol{\mu}_{\mathbf{X}_k}$						
skewness	0.4513	0.3886	0.1150	0.4504	0.0422	0.0317
kurtosis	0.0171	0.0375	0.6161	0.4693	0.4039	0.8658
$\mathbf{y} - \mathcal{X}' oldsymbol{eta}_k$						
skewness	0.6053	0.6263	0.5742	0.2526	0.6386	0.0212
kurtosis	0.0369	0.3261	0.9268	0.3922	0.1747	0.7168

5. Conclusions

The proposed multivariate seemingly unrelated Gaussian linear cluster-weighted 811 models can account for heterogeneous regression data with multivariate corre-812 lated responses, each one depending on its own set of covariates. This latter 813 feature represents the main novelty of the models proposed here in reference 814 with the ones introduced by Dang et al. (2017), thus leading to a more flex-815 ible modelling of data in applications where prior information concerning the 816 absence of certain covariates from the linear term employed in the prediction 817 of a certain response has to be conveyed into the model, and different covari-818 ates are expected to be relevant in the prediction of different responses. The distribution of the covariates is also explicitly incorporated in the model for-820 mulation. The resulting approach encompasses the models introduced by Dang 821 et al. (2017) as well as other Gaussian mixture-based linear regression models 822 with random covariates. Details about identifiability, ML estimation and model 823 selection have been provided. Furthermore, models with a reduced number of variance-covariance parameters have been specified. The comparisons among 825 some cluster-weighted models and clusterwise linear regression models based on 826 the analyses of the tuna and RtI datasets have highlighted the effectiveness of 827 the proposed models in detecting the presence of unobserved heterogeneity; such models have been proved to be useful also to establish the relevance of a multi-829 variate regression analysis and the inadequacy of the assignment independence 830 assumption in both applications. From the Monte Carlo studies it appears that 831 including irrelevant regressors in a cluster-weighted model can lead to a wrong 832 choice of the number of components and a sub-optimal reconstruction of the true classification of the sample observations, especially when the components 834 are not well-separated. The approach introduced here is able to avoid some 835 drawbacks due to the presence of irrelevant regressors in a multivariate Gaus-836 837 sian linear cluster-weighted model. This happens because the proposed models are multivariate Gaussian linear cluster-weighted models in which some regression coefficients are set a priori equal to zero. Thus, the proposed approach also 839

represents a framework for multivariate linear cluster-weighted analysis under such constraints.

As far as the development of inferential methods for the parameters of the proposed models is concerned, an assessment of the sample variability of the pa-843 rameter estimates is required. Since the ECM algorithm does not automatically 844 produce any estimate of the covariance matrix of the ML estimator, additional 845 computations are necessary. To this end, several approaches commonly employed under finite mixture models could be exploited (see, e.g., McLachlan and Peel, 2000). For example, estimates of the asymptotic covariance matrix 848 of the ML estimator can be computed through an approach which is based on 849 the gradient vector and the second-order derivative matrix of the incomplete 850 data log-likelihood, and makes also use of a sandwich estimator. This approach 85 has been successfully applied to Gaussian mixture models (Boldea and Magnus, 2009), t mixture models (Wang and Lin, 2016), clusterwise Gaussian linear re-853 gression models (Galimberti et al., 2021) and Gaussian linear cluster-weighted 854 models (Soffritti, 2021). In addition, given the critical role played by the initial-855 isation in any ECM algorithm, further investigation might be needed in order 856 to confirm the encouraging results described in Section 3. In particular, this ad-857 ditional investigation should focus on the performance of the proposed strategy 858 in presence of high dimensional data. 859

Another crucial aspect associated with the adoption of the proposed models in practical applications is the assessment of their adequacy. For finite mixtures of linear regression models with a univariate response and fixed, concomitant or random covariates, Ingrassia and Punzo (2020) have recently introduced some indices able to measure the association between the response variable and the latent groups, the model goodness-of-fit, and the proportion of the total variation in the response which remains unexplained by the fitted model. Local and overall coefficients of determination have also been described. After suitable modifications, those indices could also be employed to assess the adequacy of the multivariate cluster-weighted models based on seemingly underlated linear regression illustrated here.

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Multivariate seemingly unrelated linear cluster-weighted analyses based on 871 the proposed models implicitly require that the researcher has prior information 872 on the specific covariates that have to be included in the linear term employed in the prediction of each response in the model. In practical applications in which 874 the choice of the regressors to be used for different responses is questionable, the 875 relevant regressors for each response can be detected through strategies (e.g., 876 stepwise techniques, genetic algorithms) that allow to perform variable selec-877 tion in a multivariate regression framework. To this end, the optimal model for 878 the given dataset should be determined from a model class which also includes 879 the cluster-weighted models based on seemingly unrelated linear regression il-880 lustrated in this paper. 881

Finally, it is worth noting that the new multivariate seemingly unrelated 882 cluster-weighted models described here have been specified under the following assumptions: i) the joint conditional distribution of the P covariates given 884 the group Ω_k is Gaussian $\forall k$; ii) the joint conditional distribution of the D 885 responses given the covariates and the group Ω_k is Gaussian $\forall k; iii)$ the con-886 ditional expected value of the D responses given the covariates and the group 887 Ω_k is a linear transformation of the covariates $\forall k$. These assumptions could 888 be relaxed by resorting to the approaches developed by Punzo (2014), Punzo 889 and McNicholas (2017), Gallaugher et al. (2021) or Sahin and Czado (2021) 890 so as to obtain multivariate seemingly unrelated cluster-weighted models which 891 could be more effectively employed in the analysis of real datasets composed of unknown clusters of observations characterised by skewed distributions, outliers 893 or non-linear relationships. 894

Appendix A. Derivation of $\hat{eta}_k^{*(h+1)}$ and $\hat{\Sigma}_{\mathrm{Y}_k}^{(h+1)}$

The CM-steps to update the estimates of the model parameters β_k^* and $\Sigma_{\mathbf{Y}_k}$ at the (h+1)th iteration in the ECM algorithm, as illustrated in equations (13)

and (14), can be obtained as follows.

$$\frac{\partial}{\partial \boldsymbol{\beta}_{k}^{*'}} Q(\boldsymbol{\psi}|\boldsymbol{\psi}^{(h)}) = \frac{\partial}{\partial \boldsymbol{\beta}_{k}^{*'}} \sum_{i=1}^{I} \sum_{k=1}^{K} \hat{\tau}_{ik}^{(h)} Q_{2}(\boldsymbol{\beta}_{k}^{*}, \boldsymbol{\Sigma}_{\mathbf{Y}_{k}}|\boldsymbol{\psi}^{(h)})$$

$$= \frac{\partial}{\partial \boldsymbol{\beta}_{k}^{*'}} \left[-\frac{1}{2} \sum_{i=1}^{I} \hat{\tau}_{ik}^{(h)} (\mathbf{y}_{i} - \boldsymbol{\mathcal{X}}_{i}^{\prime} \boldsymbol{\beta}_{k}^{*})^{\prime} \boldsymbol{\Sigma}_{\mathbf{Y}_{k}}^{(h)(-1)} (\mathbf{y}_{i} - \boldsymbol{\mathcal{X}}_{i}^{\prime} \boldsymbol{\beta}_{k}^{*}) \right]$$

$$= -\frac{1}{2} \sum_{i=1}^{I} \hat{\tau}_{ik}^{(h)} \frac{\partial}{\partial \boldsymbol{\beta}_{k}^{*'}} \left(-2\mathbf{y}_{i}^{\prime} \boldsymbol{\Sigma}_{\mathbf{Y}_{k}}^{(h)(-1)} \boldsymbol{\mathcal{X}}_{i}^{\prime} \boldsymbol{\beta}_{k}^{*} + \boldsymbol{\beta}_{k}^{*'} \boldsymbol{\mathcal{X}}_{i}^{\prime} \boldsymbol{\Sigma}_{\mathbf{Y}_{k}}^{(h)(-1)} \boldsymbol{\mathcal{X}}_{i}^{\prime} \boldsymbol{\beta}_{k}^{*} \right)$$

$$= -\frac{1}{2} \sum_{i=1}^{I} \hat{\tau}_{ik}^{(h)} \left(-2\mathbf{y}_{i}^{\prime} \boldsymbol{\Sigma}_{\mathbf{Y}_{k}}^{(h)(-1)} \boldsymbol{\mathcal{X}}_{i}^{\prime} + 2\boldsymbol{\beta}_{k}^{*'} \boldsymbol{\mathcal{X}}_{i} \boldsymbol{\Sigma}_{\mathbf{Y}_{k}}^{(h)(-1)} \boldsymbol{\mathcal{X}}_{i}^{\prime} \right)$$

$$= \sum_{i=1}^{I} \hat{\tau}_{ik}^{(h)} \mathbf{y}_{i}^{\prime} \boldsymbol{\Sigma}_{\mathbf{Y}_{k}}^{(h)(-1)} \boldsymbol{\mathcal{X}}_{i}^{\prime} - \sum_{i=1}^{I} \hat{\tau}_{ik}^{(h)} \boldsymbol{\beta}_{k}^{*'} \boldsymbol{\mathcal{X}}_{i} \boldsymbol{\Sigma}_{\mathbf{Y}_{k}}^{(h)(-1)} \boldsymbol{\mathcal{X}}_{i}^{\prime} . \tag{A.1}$$

Setting (A.1) equal to the null vector, $\Sigma_{\mathbf{Y}_k}^{(h)}$ equal to $\hat{\Sigma}_{\mathbf{Y}_k}^{(h)}$ and solving the so obtained system with respect to $\boldsymbol{\beta}_k^*$ leads to the solution reported in equation (13).

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$$\frac{\partial}{\partial \mathbf{\Sigma}_{\mathbf{Y}_{k}}^{-1}} Q(\boldsymbol{\psi}|\boldsymbol{\psi}^{(h)}) = \frac{\partial}{\partial \mathbf{\Sigma}_{\mathbf{Y}_{k}}^{-1}} \sum_{i=1}^{I} \sum_{k=1}^{K} \hat{\tau}_{ik}^{(h)} Q_{2}(\boldsymbol{\beta}_{k}^{*}, \mathbf{\Sigma}_{\mathbf{Y}_{k}}|\boldsymbol{\psi}^{(h)})$$
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$$= \frac{\partial}{\partial \mathbf{\Sigma}_{\mathbf{Y}_{k}}^{-1}} \left\{ \frac{1}{2} \sum_{i=1}^{I} \hat{\tau}_{ik}^{(h)} \left[\ln |\mathbf{\Sigma}_{\mathbf{Y}_{k}}^{-1}| \right. \right.$$
908
$$- \left(\mathbf{y}_{i} - \mathcal{X}_{i}' \boldsymbol{\beta}_{k}^{*(h+1)} \right)' \mathbf{\Sigma}_{\mathbf{Y}_{k}}^{-1} \left(\mathbf{y}_{i} - \mathcal{X}_{i}' \boldsymbol{\beta}_{k}^{*(h+1)} \right) \right] \right\}$$
909
$$= \frac{1}{2} \sum_{i=1}^{I} \hat{\tau}_{ik}^{(h)} \frac{\partial}{\partial \mathbf{\Sigma}_{\mathbf{Y}_{k}}^{-1}} \left[\ln |\mathbf{\Sigma}_{\mathbf{Y}_{k}}^{-1}| \right.$$
910
$$- tr \left(\mathbf{\Sigma}_{\mathbf{Y}_{k}}^{-1} (\mathbf{y}_{i} - \mathcal{X}_{i}' \boldsymbol{\beta}_{k}^{*(h+1)}) (\mathbf{y}_{i} - \mathcal{X}_{i}' \boldsymbol{\beta}_{k}^{*(h+1)})' \right) \right]$$
911
$$= \frac{1}{2} \left[\sum_{i=1}^{I} \hat{\tau}_{ik}^{(h)} \mathbf{\Sigma}_{\mathbf{Y}_{k}} - \sum_{i=1}^{I} \hat{\tau}_{ik}^{(h)} (\mathbf{y}_{i} - \mathcal{X}_{i}' \boldsymbol{\beta}_{k}^{*(h+1)}) (\mathbf{y}_{i} - \mathcal{X}_{i}' \boldsymbol{\beta}_{k}^{*(h+1)}) (\mathbf{y}_{i} - \mathcal{X}_{i}' \boldsymbol{\beta}_{k}^{*(h+1)})^{2} \right]$$

where the second and third equalities are obtained using properties of trace and

₉₁₃ transpose and differentiation rules of functions of matrices. Setting (A.2) equal

to the null matrix, $oldsymbol{eta}_k^{*(h+1)}$ equal to $\hat{oldsymbol{eta}}_k^{*(h+1)}$ and solving the resulting system

with respect to $\Sigma_{\mathbf{Y}_k}$ gives the update in equation (14).

Appendix B. Expression of $\hat{\beta}_k^{*(h+1)}$ when $\mathbf{x}_{id} = \mathbf{x}_i \, \forall d$

Similarly to Park (1993), equation (13) can be rewritten as

$$\hat{\boldsymbol{\beta}}_{k}^{*} = \left\{ \mathcal{X} \left[\operatorname{diag} \left(\hat{\boldsymbol{\tau}}_{k} \right) \otimes \hat{\boldsymbol{\Sigma}}_{\mathbf{Y}_{k}}^{-1} \right] \mathcal{X}' \right\}^{-1} \mathcal{X} \left[\operatorname{diag} \left(\hat{\boldsymbol{\tau}}_{k} \right) \otimes \hat{\boldsymbol{\Sigma}}_{\mathbf{Y}_{k}}^{-1} \right] \mathbf{y}, \tag{B.1}$$

where the superscripts (h) and (h+1) have been dropped to ease notation, $\mathcal{X} = [\mathcal{X}_1, \mathcal{X}_2, \dots, \mathcal{X}_I]$ is a $(P^* + D) \times (D \cdot I)$ matrix, diag $(\hat{\tau}_k)$ is a diagonal matrix whose diagonal elements are the values $\hat{\tau}_{ik}$ $(i = 1, \dots, I)$ and $\mathbf{y} = (\mathbf{y}'_1, \mathbf{y}'_2, \dots, \mathbf{y}'_I)'$. Consider now the vectors $\mathbf{v}_d = (y_{1d}, y_{2d}, \dots, y_{Id})'$, containing the values of the dth response on the I observations $(d = 1, \dots, D)$, and
the vector $\mathbf{v} = (\mathbf{v}'_1, \dots, \mathbf{v}'_D)'$. It is evident that \mathbf{v} and \mathbf{y} contain the same values
but in a different order. As shown in Park (1993), by exchanging the rows of
the identity matrix of order $D \cdot I$, it is possible to define a matrix \mathbf{L} such that

$$\mathbf{L}\mathbf{L}' = \mathbf{L}'\mathbf{L} = \mathbf{I}_{D \cdot I}$$

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$$\mathbf{L}\mathbf{y} = \mathbf{v}.$$

Matrix **L** can also be used to reorder the columns of \mathcal{X} and the rows and columns of $\left[\operatorname{diag}\left(\hat{\boldsymbol{\tau}}_{k}\right)\otimes\hat{\boldsymbol{\Sigma}}_{\mathbf{Y}_{k}}^{-1}\right]$. Namely,

$$\mathcal{X}\mathbf{L}' = \mathcal{Z} = egin{bmatrix} \mathbf{Z}_1 & \mathbf{0}_{(P_1+1) imes I} & \dots & \mathbf{0}_{(P_1+1) imes I} \ \mathbf{0}_{(P_2+1) imes I} & \mathbf{Z}_2 & \dots & \mathbf{0}_{(P_2+1) imes I} \ dots & dots & dots \ \mathbf{0}_{(P_D+1) imes I} & \mathbf{0}_{(P_D+1) imes I} & \dots & \mathbf{Z}_D \end{bmatrix},$$

where $\mathbf{Z}_d = [\mathbf{x}_{1d}^*, \mathbf{x}_{2d}^*, \cdots, \mathbf{x}_{Id}^*]$ is a $(P_d + 1) \times I$ matrix $(d = 1, \dots, D)$, and

$$\mathbf{L}\left[\operatorname{diag}\left(\hat{\boldsymbol{\tau}}_{k}\right)\otimes\hat{\boldsymbol{\Sigma}}_{\mathbf{Y}_{k}}^{-1}\right]\mathbf{L}'=\left[\hat{\boldsymbol{\Sigma}}_{\mathbf{Y}_{k}}^{-1}\otimes\operatorname{diag}\left(\hat{\boldsymbol{\tau}}_{k}\right)\right].$$

Thus, an equivalent expression for $\hat{\boldsymbol{\beta}}_k^*$ is given by

$$\hat{\boldsymbol{\beta}}_{k}^{*} = \left\{ \mathcal{X} \mathbf{L}' \mathbf{L} \left[\operatorname{diag} \left(\hat{\boldsymbol{\tau}}_{k} \right) \otimes \hat{\boldsymbol{\Sigma}}_{\mathbf{Y}_{k}}^{-1} \right] \mathbf{L}' \mathbf{L} \mathcal{X}' \right\}^{-1} \mathcal{X} \mathbf{L}' \mathbf{L} \left[\operatorname{diag} \left(\hat{\boldsymbol{\tau}}_{k} \right) \otimes \hat{\boldsymbol{\Sigma}}_{\mathbf{Y}_{k}}^{-1} \right] \mathbf{L}' \mathbf{L} \mathbf{y} \right]$$

$$= \left\{ \mathcal{Z} \left[\hat{\boldsymbol{\Sigma}}_{\mathbf{Y}_{k}}^{-1} \otimes \operatorname{diag} \left(\hat{\boldsymbol{\tau}}_{k} \right) \right] \mathcal{Z}' \right\}^{-1} \mathcal{Z} \left[\hat{\boldsymbol{\Sigma}}_{\mathbf{Y}_{k}}^{-1} \otimes \operatorname{diag} \left(\hat{\boldsymbol{\tau}}_{k} \right) \right] \mathbf{v}.$$
(B.2)

If $\mathbf{x}_{id}=\mathbf{x}_i \ \forall d$, then $\mathbf{Z}_d=[\mathbf{x}_1^*,\mathbf{x}_2^*,\cdots,\mathbf{x}_I^*]=\mathbf{Z} \ \forall d$ and

$$\mathcal{Z} = \mathbf{I}_D \otimes \mathbf{Z}. \tag{B.3}$$

By exploiting equation (B.3) and the properties of the Kronecker product (see,
e.g., Magnus and Neudecker, 1988), equation (B.2) can be simplified as follows:

$$\hat{\boldsymbol{\beta}}_{k}^{*} = \left\{ \mathbf{I}_{D} \otimes \left[\mathbf{Z} \operatorname{diag} \left(\hat{\boldsymbol{\tau}}_{k} \right) \mathbf{Z}' \right]^{-1} \mathbf{Z} \operatorname{diag} \left(\hat{\boldsymbol{\tau}}_{k} \right) \right\} \mathbf{v}. \tag{B.4}$$

Firstly, note that equation (B.4) does not depend on $\hat{\Sigma}_{\mathbf{Y}_k}$. Furthermore, the matrix $\left\{\mathbf{I}_D \otimes \left[\mathbf{Z} \operatorname{diag}(\hat{\boldsymbol{\tau}}_k) \mathbf{Z}'\right]^{-1} \mathbf{Z} \operatorname{diag}(\hat{\boldsymbol{\tau}}_k)\right\}$ has a block-diagonal structure.

By coupling it with the structure of the vector \mathbf{v} , the following expression for the vector $\hat{\boldsymbol{\beta}}_{kd}^*$ containing the estimated coefficients associated with the dth response in the kth group can be obtained:

$$\hat{\boldsymbol{\beta}}_{kd}^{*} = \left[\mathbf{Z} \operatorname{diag}\left(\hat{\boldsymbol{\tau}}_{k}\right) \mathbf{Z}'\right]^{-1} \mathbf{Z} \operatorname{diag}\left(\hat{\boldsymbol{\tau}}_{k}\right) \mathbf{v}_{d}$$

$$= \left(\sum_{i=1}^{I} \hat{\tau}_{ik} \mathbf{x}_{i}^{*} \mathbf{x}_{i}^{*}'\right)^{-1} \left(\sum_{i=1}^{I} \hat{\tau}_{ik} \mathbf{x}_{i}^{*} y_{id}\right), \quad d = 1, \dots, D. \quad (B.5)$$

Apart from differences related to notation, it can be noticed that equation (B.5) coincides with the dth row of the matrix defined in equation (8) in Dang et al. (2017).

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