CorReg: model-based pretreatment for regression with correlated variables. Application in steel industry

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Abstract. Linear regression outcomes are known to be damaged by correlated covariates. However many modern datasets are expected to convey more and more highly correlated covariates. We propose to explicitly model the correlations by a family of linear regressions between the covariates. Marginalization allows then to obtain a parsimonious correlation-free regression model, easily understandable and from which it is then possible to perform standard linear estimation methods including variables selection procedures for instance. The structure of correlations is found with an MCMC algorithm. An R package (Correlation available on the CRAN implements this new method which will be illustrated on both simulated datasets and real-life datasets from steel industry where correlated variables are frequent.

Keywords. Regression, correlations, industry, variable selection, generative models

1 Introduction

When one wants to explain a phenomenon based on some covariates, the first statistical method tried frequently is the linear regression. It provides a predictive model with a good interpretability even for non-statistician and is simple to learn. Therefore, linear regression is used in nearly all the fields where statistics are made, from industry (ballistic models to calibrate the process) to sociology (predicting some numerical properties of a population). Linear regression is a very classic situation but sometimes it has to face an also classical problem: the variance of the estimators, even whithout bias from modelisation. We note the linear regression model:

$$Y_{|X} = X\beta + \varepsilon \tag{1}$$

where X is the $n \times p$ matrix of the explicative variables, Y the $n \times 1$ response vector and $\varepsilon \sim \mathcal{N}(0, \sigma_Y^2 I_n)$ the noise of the regression, with I_n the n-sized identity matrix and $\sigma_Y \in \mathcal{R}$. The $p \times 1$ vector β is the vector of the coefficients of the regression, estimated with Ordinary Least Squares (OLS). As shown in section 2.1, estimation of the parameters requires the inversion of a matrix which will be ill-conditioned or even singular if some covariates depend linearly from each other. Variance of $\hat{\beta}$ increases based on two aspects:

- The dimension p (number of covariates) of the model : the more covariates you have the greater variance you get.
- The correlations within the covariates : strongly correlated covariates give bad-conditioning and increase variance of the estimators .

With the rise of informatic, datasets contains more and more covariates and thus more and more useless covariates. So dimension reduction becomes a necessity. Moreover, when you use more covariates, you increase the chance to have correlated ones. It will be illustrated with examples from steel industry in the last section of this paper (many parameters of the whole process without any a priori) highly correlated (physical laws, process rules, etc). In such a context, variance of the estimators can lead to arbitrary results or even no results at all. Prediction and interpretation are both

strongly needed, depending on the context. For instance, in an industrial context interpretation could be favored to improve the process instead of only predict defects.

Because OLS is the minimum-variance unbiased estimator, penalized methods try to reduce the variance introducing some bias to improve the bias-variance trade-off and get better prediction. Moreover, real datasets imply many irrelevant variables so we have to use variable selection methods.

In the following we note classical norms: $\|\beta\|_2^2 = \sum_{i=1}^p (\beta_i)^2$ and $\|\beta\|_1 = \sum_{i=1}^p |\beta_i|$.

Ridge regression[13] proposes a biased estimator that can be written in terms of a parametric L_2 penalty:

 $\hat{\beta} = \operatorname{argmin} \left\{ \| Y - X\beta \|_2^2 \right\} \text{ subject to } \| \beta \|_2^2 \le k$ (2)

But Ridge regression do not aim at selecting covariates because coefficients tends to 0 but don't reach 0. So it gives difficult interpretations for large values of p.

The Least Absolute Shrinkage and Selection Operator (LASSO [17]) consists in a shrinkage of the regression coefficients based on a λ parametric L_1 penalty.

$$\hat{\beta} = \operatorname{argmin} \left\{ \| Y - X\beta \|_{2}^{2} \right\} \text{ subject to } \| \beta \|_{1} \leq \lambda$$
 (3)

The Least Angle Regression (LAR [4]) Algorithm offers a very efficient way to obtain the whole LASSO path and is very attractive. It requires only the same order of magnitude of computational effort as OLS applied to the full set of covariates. And it really selects covariates with coefficients set exactly to 0. But LASSO also faces consistency problems [23] when confronted with correlated covariates. Another limitation of the LASSO is that it preserves at most n predictors (troublesome when in high dimension). Some recent variants of the LASSO does exist for the choice of the penalization coefficient like the adaptative LASSO [24] or the random LASSO [19].

Elastic net[25] is a method developed to be a compromise between Ridge regression and the LASSO. Elastic net can be written:

$$\hat{\beta} = (1 + \lambda_2) \operatorname{argmin} \left\{ \| Y - X\beta \|_2^2 \right\}, \text{ subject to } (1 - \alpha) \| \beta \|_1 + \alpha \| \beta \|_2^2 \le t \text{ for some } t$$
 (4)

where $\alpha = \frac{\lambda_2}{(\lambda_1 + \lambda_2)}$. But it is based on the grouping effect and if the dataset contains two identical variables they will obtain the same coefficient whereas LASSO will choose between one of them and will then obtain same predictions with a more parsimonious model.

Another way of reducing the dimension is to consider clusters of variables with the same coefficients, like the Octogonal Shrinkage and Clustering Algorithm for Regression (OSCAR [1]). The CLusterwise Effect REgression (CLERE [20]) describes the β_j no longer as fixed effect parameters but as unobserved independent random variables whith grouped β_j following a Gaussian Mixture distribution.

The idea is to hope that the model have a small number of groups of covariates and that the mixture will have few enough components to have a number of parameters to estimate significantly lower than p. In such a case, it improves interpretability and ability to yeld reliable prediction with a smaller variance on $\hat{\beta}$. But it requires to suppose having many covariates with the same level of effect on the response variable and seems to stay less efficient in prediction than elastic net. Spike and Slab variable selection [10] also relies on Gaussian mixture (the spike and the slab) hypothesis for the β_j and gives a subset of covariates (not grouped) on which to compute OLS but has no specific protection against correlations issues.

Where some try to reduce the dimension and then have to suppose having no correlations between the remaining covariates, we propose to focus on the correlations, giving a model with orthogonal covariates and an explicit structure between covariates. In fact we search the greatest set of orthogonal covariates to keep the maximum information but with an orthogonality constraint. This can be viewed as a pretreatment on the dataset allowing to use then other dimension reduction tools without suffering from correlations.

Our work is based on the assumption that if we know explicitly the source of correlations, we could use this knowledge to avoid the problem. The structure obtained gives a system of linear regression that can be viewed as a recursive Simultaneous Equation Model (SEM)[3]. Such a system is easy to interpret but estimation don't take advantage of the explicit structure [18] when the structure is straight forward (recursive SEM). Other methods take into account correlations but they only consider covariances between residuals SUR (Seemingly Unrelated Regression [21]) or endogenous variables like SPRING (Structured selection of Primordial Relationships IN the General linear model [2]).

We make the choice to distinguish the response variable from the other endogenous variables (that are on the left of a regression). Thus we have one regression on our response variable and a system of sub-regressions (without the response variable). Then we consider correlations between the explicative covariates of the main regression, not between the residuals. The structure is supposed to be the source of the correlations and allows us to define a marginal regression model based on a reduced set of independent covariates without significant information loss. So we may obtain two kinds of zeros in the marginal model: coerced zeros due to correlations (redundant information) and estimated ones with classical variable selection methods applied on remaining variables. This two kinds of zero won't be interpreted in the same way and thus consistency issues don't mean interpretation issues any more. So we dodge the drawbacks of both grouping effect and variable selection.

SEM are often used in social sciences and economy where a structure is supposed "by hand" but here we want to find it automatically. Graphical LASSO [6] offers a method to obtain a structure based on the precision matrix (inverse of the variance-covariance matrix). It consists in a selection in the precision matrix, setting some covariances to zero. But the resulting matrix is symmetric and we need an oriented structure for our SEM. So we developped an MCMC algorithm to find the structure (R package Correction of CRAN). The structure is based on a full generative model on X to be able to compare likelihood-based criterion in the MCMC.

This paper will first present the marginal regression model and its properties before describing in Section 3 the random walk used to find the structure. We will then look at some numerical results on simulated (Section 4) and real industrial datasets (Section 5) before concluding and giving some perspectives in the last part.

$\mathbf{2}$ Model to select decorrelated covariates

The decorrelation problem in regression 2.1

For a model defined by

$$Y_{|X} = X\beta + \varepsilon \tag{5}$$

where X is the $n \times p$ matrix of the explicative variables, Y the response vector and $\varepsilon \sim \mathcal{N}(0, \sigma_V^2 I_n)$ with I_n the n-sized identity matrix, we have the following Ordinary Least Squares (OLS) estimators [15]:

$$\hat{\beta} = (X'X)^{-1}X'Y \tag{6}$$

With variance

$$\operatorname{Var}(\hat{\beta}) = \sigma_Y^2 (X'X)^{-1} \tag{7}$$

The Mean Squared Error (MSE) on $\hat{\beta}$ is:

$$MSE(\hat{\beta}|\beta, X) = Bias^{2} + Tr(Var(\hat{\beta}))$$

$$= 0 + \sigma_{Y}^{2} Tr((X'X)^{-1})$$
(8)

$$= 0 + \sigma_Y^2 \operatorname{Tr}((X'X)^{-1})$$
 (9)

and when correlations between covariates are strong, the matrix to invert is ill-conditioned, the variance explodes and so does the MSE. Another problem is that matricial inversion requires $n \geq p$. When it is not the case, a frequently used alternative is the Moore-Penrose generalized inverse [11]. In the followings, matrices inverses when n < p will refer to this generalized inverse. Thus OLS can obtain some results even with n < p (see section 4).

2.2 Modelisation of the correlations

Let now $X \in \mathbb{R}^{n \times p}$ be a set of p correlated covariates. We propose to explicitly define a family of p_2 internal regressions between covariates whith I_2 the set of indices of endogenous variables in X (explained ones) and $I_1 = \{I_1^1, \ldots, I_1^p\}$ the set of the sets of indices of exogenous covariates (explaining ones) with $\forall j \notin I_2, I_1^j = \emptyset$. Then we have an explicit structure $S = (I_1, I_2, p_1, p_2)$ where $p_1 = (p_1^1, \ldots, p_1^{p_2})$ is the vector of the number of covariates in each internal regression. Thus we have $p_2 = |I_2|$ and $p_1^j = |I_1^j|$ where |.| represents the cardinal of an ensemble.

In the following, we note X^j the j^{th} column of a matrix X. For lighter notation we define $X_2 = X^{I_2}$ the matrix of the endogenous covariates and $X_1 = X \setminus X_2$ the matrix of the remaining exogenous covariates.

We define a family of p_2 regressions within correlated covariates in X:

$$\forall j \in I_2 : X_{|X_1,S}^j = X_1 B_1^j + \varepsilon_j \text{ with } \varepsilon_j \sim (0, \sigma_j^2 I_n)$$
(10)

where $B_1^j \in \mathcal{R}^{(p-p_2)}$ are the vectors of the regression coefficients between the covariates (containing some zeros according to I_1^j). It remains $p-p_2$ independent exogenous variables: $\forall j \notin I_2: X_{|S|}^j \sim f(\theta_j)$

We make the hypothesis of the uncrossing rule $I_1 \cap I_2 = \emptyset$ *i.e.* endogenous variables don't explain other covariates, thus we have a partition $X = [X_1, X_2]$.

This generative model is conditional to S, the discrete structure model that is identifiable because we can't permute some regressions in (10) and obtain the same joint distribution P(X,Y), the residuals (ε_j) would not stay Gaussian (see the appendices). These residuals are in the following supposed independent but one can suppose dependencies between them and then use appropriate tools to estimate them and the B_1^j like SUR with Feasible Generalized Least Squares (FGLS) by Zellner [21] or SPRING [2] as mentioned in the introduction.

If there are exact regressions $(\sigma_j^2 = 0)$ in (10), the structure won't be identifiable. But it only means that several structure will have the same likelihood and they will have the same interpretation. So it's not really a problem. Moreover, when an exact sub-regression is found, we can delete one of the implied variables without any loss of information and the structure will define a list of variable from which to delete.

2.3 Marginal regression model with decorrelated covariates

Let $Y \in \mathbb{R}^n$ be a response variable we want to explain with X. We note that (11) and (10) also give by simple integration on X_2 a regression model on Y depending only on uncorrelated covariates X_1 :

$$Y_{|S,X} = X\beta + \varepsilon_Y = X_1\beta_1 + X_2\beta_2 + \varepsilon_Y \text{ whith } \varepsilon_Y \sim \mathcal{N}(0, \sigma_Y^2 I_n);$$
 (11)

where $\beta = (\beta_1, \beta_2) \in \mathcal{R}^p$ is the vector of the regression coefficients and I_n the identity matrix. Knowing (10) we then obtain a marginal regression model:

$$Y_{|X_1} = X_1(\beta_1 + \sum_{j \in I_2} B_1^j \beta_j) + \sum_{j \in I_2} \varepsilon_j \beta_j + \varepsilon_Y$$
(12)

$$= X_1 \alpha_1 + X_2 \alpha_2 + \varepsilon_{\alpha} \text{ where } \alpha_2 = 0.$$
 (13)

We note that it simply is a linear regression on some of the original covariates so we only made a pretreatment on the dataset by setting some coefficients to 0 because of correlations. These 0 won't be interpreted as independence with the response variable but as redundant information due to correlations between the covariates. It is a variable pre-selection independent of the response Y. As a pretreatment, it allows usage of any method in a seconde time to estimate α .

2.4 Estimator and properties

The marginal regression model simply is

$$Y_{|X_1} = X_1 \alpha_1 + \varepsilon_\alpha \tag{14}$$

So we have the unbiased estimator of α :

$$\hat{\alpha}_1 = (X_1' X_1)^{-1} X_1' Y \text{ and } \hat{\alpha}_2 = 0$$
 (15)

We see in (12) that it gives an unbiased estimation of Y and α but in terms of β this estimator is biased:

$$E[\hat{\alpha}_1|X_1] = \beta_1 + \sum_{j \in I_2} B_1^j \beta_j \text{ and } E[\hat{\alpha}_2|X_1] = 0$$
 (16)

with variance:

$$\operatorname{Var}[\hat{\alpha}_1|X_1] = (\sigma_Y^2 + \sum_{j \in I_2} \sigma_j^2 \beta_j^2)(X_1' X_1)^{-1} \text{ and } \operatorname{Var}[\hat{\alpha}_2|X_1] = 0$$
(17)

We see that the variance is reduced compared to OLS described in equation (7)(no correlations and smaller matrix give better conditioning) for small values of σ_j i.e. strong correlations. Moreover, the explicit structure is parsimonious and simply consists in linear regressions and thus is easily understood by non statistician, allowing them to have a better knowledge of the phenomenon inside the dataset. Expert knowledge can even be added to the structure. This new model is reduced even without variable selection and is just a linear regression so every method for variable selection in linear regression can be used then, like LASSO or even stepwise [16]. Hence we hope to obtain a parsimonious model with two kinds of zeros: those from decorrelating step and those from selection step, with different meanings. A last Ordinary Least Square step is added to improve estimation because penalisation is made to select variables but also shrinks remaining coefficients. This last step allows to keep the benefits of shrinkage (variable selection) without any impact on remaining coefficients (see [22]).

To better illustrate the bias-variance compromise, we look at a simple example with p=3 variables. X_1 is composed by two independent scaled Gaussian $\mathcal{N}(0,1)$, $X_3=\frac{1}{2}x_1+\frac{1}{2}x_2+\varepsilon_3$ where $\varepsilon_3\sim\mathcal{N}(0,\sigma_3^2)$. We also have $\beta=(\frac{1}{3},\frac{1}{3},\frac{1}{3})$ and $\sigma_Y=10$. Then we observe the theoretical Mean Squared Error (MSE) of the estimator of both OLS and Correspondence: $MSE(\hat{\beta}|\beta,X)=Bias^2+\mathrm{Tr}(\mathrm{Var}(\hat{\beta}))$ for several values of σ_3 (strength of the sub-regression) and n. Figures 1 to 3 show the theoretical MSE evolution with the strength of the sub-regression:

$$1 - \mathcal{R}^2 = \frac{\operatorname{Var}(\varepsilon_3)}{\operatorname{Var}(x_3)} = \frac{\sigma_3^2}{\sigma_3^2 + \frac{1}{2}}$$
(18)

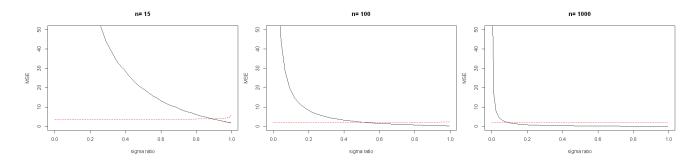


Figure 1: For n=15. Dotted: Figure 2: For n=100. Dotted: Figure 3: For n=1000. Dotted: Correg, plain: OLS Correg, plain: OLS Correg, plain: OLS.

It is clear in Figures 1 to 3 that the reduced model is more robust than OLS. And even when subregression are week (small value of $1 - \mathcal{R}^2$) it does not explode. We also see that the error implied by strong correlations shrinks whit the rise of n. $MSE_{OLS} = 0 + \text{Tr}(\text{Var}(\hat{\beta})) = \text{Tr}(\text{Var}(\hat{\beta}_1)) + \text{Tr}(\text{Var}(\hat{\beta}_2))$. both terms are multiplied by σ_Y^2 and so is $\text{Tr}(\text{Var}(\hat{\alpha}_1))$. Thus, as $\text{Var}(\hat{\alpha}_1) = 0$ when σ_Y^2 rises it increases the advantage of Corresponding of Corresponding to the importance of dimension reduction when the model has a strong noise (very usual case on real datasets where true model is not even exactly linear).

2.5 Advantages for interpretation

Grouping effect is the fact that correlated covariates get similar coefficient and are selected together. It's the case with elasticnet [25] for example. If $X_2 = X_1 + e$ and we have the grouping effect, we will obtain a model like $\hat{Y} = \hat{\beta}X_1 + \hat{\beta}X_2$. Then, if you try to modify the response value, you will modify one of the covariates and both will change so you won't get expected results.

Nothing constrains us to give only one equation. If you have the following model: $Y = \beta X_1 + \beta X_2$ and $X_1 = X_2 + e$. Then you have more information and are able to decide better actions. With such a model, grouping effect is no more useful because when saying $Y = 2\beta X_1$ and $X_2 = X_1 + e$ with the information that X_2 has been removed locally because of its correlation with X_1 , you don't get misleading interpretations anymore. In fact, we have two kinds of zeros (those from selection and those from correlations) and we know for each zero which kind is it. So it is possible to combine the advantages of grouping effect and selection just giving several equations. Each equation here is very simple (only linear regressions) so you don't really increase complexity of the model. Moreover, the uncrossing constraint $(I_1 \cap I_2 = \emptyset)$ guarantee to keep a simple structure easily interpretable (no cycles and no chain-effect) and straightforward readable.

3 Estimating structure of sub-regressions with a Markov chain

3.1 Detailed generative model

To be able to compare structures with probabilistic criterions, we need a full generative model. We suppose that variables in X_1 follow Gaussian mixtures.

$$\forall j \notin I_2 : X_{|S}^j \sim f(\theta_j) = \mathcal{GM}(\pi_j; \mu_j; \sigma_j^2) \text{ with } \pi_j, \mu_j, \sigma_j^2 \text{ vectors of size } k_j;$$
 (19)

The great flexibility [14] of such models makes our model more robust but one can use other laws if needed. Gaussian case is just a special case $(k_j = 1)$ of Gaussian mixture so it is included in our hypothesis.

Variables in X_1 are supposed to be independent. Thus if one have some hypothesis on the distribution of some variables (exponentially distributed for example) it is possible to use it without impacting the model in other ways. We now have a full generative model.

3.2 Structure comparison

To obtain the marginal regression model we need S, the linear structure between the covariates. Our full generative generative model allows us to compare structures with criterions like the Bayesian Information Criterion (BIC) which penalize the log-likelihood according to the complexity of the structure [12]. We will prefer this kind of comparison criterion instead of cross-validation that is very time-consuming and thus not friendly with combinatory problematics. We note Θ the set of the parameters of the generative model

$$P(S|X) \propto P(X|S)P(S)$$
 (20)

$$-2\log P(X|S) \approx BIC = -2\mathcal{L}(X, S, \Theta) + |\Theta|\log(n)$$
 (21)

(22)

But BIC tends to give too complex structures because we test a great range of models. Thus we choose to penalise the complexity a bit more with a hierarchical uniform a priori law $P(S) = P(I_1|p_1,I_2,p_2)P(p_1|I_2,p_2)P(I_2|p_2)P(p_2)$ instead of a simple uniform law on S. It increases penalty on complexity for $p_2 < \frac{p}{2}$ and $p_1^j < \frac{p}{2}$. Hence this constraint on \hat{p}_2 and \hat{p}_1^j is made in the MCMC in the followings. But we can imagine to use other criterions, like the RIC (Risk Inflation Criterion [5]) that choose a penalty in $\log p$ instead of $\log n$ and thus gives more parsimonious models when p is larger than n (high dimension) or any other criterion [7] thought to be better in a given context.

We will now use the following notation: $\psi(S) = \psi(X|S)$ where $\psi(X|S)$ is the chosen criterion, that will be BIC with hierarchical uniform hypothesis in our numerical results. We do not change BIC

but only P(S) so the properties are the same as classical BIC but we will obtain better results when our hypothesis is verified.

3.3 The neighbourhood

Let's define S the ensemble of feasible structures (those with $I_1 \cap I_2 = \emptyset$). For each step, starting from $S \in S$ we define a neighbourhood:

$$\mathcal{V}_{S,j} = \{S\} \cup \{S^{(i,j)} | 1 \le i \le p, i \ne j\}$$
(23)

where
$$j \sim \mathcal{U}(\{1,\ldots,p\})$$
 (24)

With $S^{(i,j)}$ defined by the following algorithm :

- if $i \notin I_i^j$ (add):
 - $-I_1^j = I_1^j \cup \{i\}$
 - $-I_1^i = \emptyset$ (explicative variables can't depend on others : column-wise relaxation)
 - $-I_1 = I_1 \setminus \{j\}$ (dependent variables can't explain others: row-wise relaxation)
- else (remove): $I_1^j = I_1^j \setminus \{i\}$

At every moment, coherence between I_1 and others parts of S can be done by $\forall 1 \leq j \leq p : p_1^j = |I_1^j|$, $I_2 = \{j | p_1^j > 0\}, \ p_2 = |I_2|$,.

3.4 The walk

3.4.1 Transition probabilities

We first make the approximation

$$P(S|X) \approx exp(\psi(S)).$$
 (25)

The algorithm follows a time-homogeneous markov chain whose transition matrix \mathcal{P} has $|\mathcal{S}|$ rows and columns (combinatory so we'll just compute the probabilities when we need them). At each step the markov chain moves with probability:

$$\forall (S, \tilde{S}) \in \mathcal{S}^2 : \mathcal{P}(S, \tilde{S}) = \sum_{j=1}^p \mathbf{1}_{\{\tilde{S} \in \mathcal{V}_{S,j}\}} \frac{\exp(-\frac{1}{2}\psi(\tilde{S}))}{\sum_{S_l \in \mathcal{V}_{S,j}} \exp(-\frac{1}{2}\psi(S_l))}$$

$$(26)$$

And S is a finite state space.

Because the walk follows a regular and thus ergodic markov chain with a finite state space, it has exactly one stationary distribution [9] and the output will be the best structure in terms of ψ which weights each candidate. Practically speaking, CORREG returns the best structure seen during the walk. Numerical results (Section 4) illustrates the efficiency of the walk when the true model really contains a linear structure or no structure at all (Table (1)) and when the structure is not linear (Table 10)).

3.4.2 Initialisation(s)

If we have some knowledge about some sub-regressions (physical models for example) we can add them in the found and/or initial structure. So the model is really expert-friendly. The initial structure can be based on a first warming algorithm taking the correlations into account. coefficients are randomly placed into I_1 , weighted by the absolute value of the correlations. We do so in the followings. Then this structure could be for example reduced by the hadamard product with the binary matrix obtained by Graphical Lasso[6] that makes selection in the precision matrix but it is time consuming.

One would rather test multiple short chains than lose time in initialisation or long chains [8]. It also helps to face local extrema. In the followings, the chain was launched with twenty initialisations.

4 Numerical results on simulated datasets

4.1 The datasets

Now we have defined the model and the way to obtain it, we can have a look on some numerical results to see if Correct keeps its promises. The Correct package has been tested on simulated datasets. Section 4.2.2 shows the results obtained in terms of \hat{S} . Sections 4.3.1 and 4.3.2 show the results obtained using only Correct, or Correct combined with other methods. Tables give both mean and standard deviation of the observed Mean Squared Errors (MSE) on a validation sample of 1000 individuals. For each simulation, p = 40, $\sigma_Y = 10$, $\sigma = 0.001$, variables X_1 follow Gaussian mixture models of $\lambda = 5$ classes which means follow Poisson's law of parameter λ and which standard deviation also is λ . The B_1^j and β_j are generated according to the same Poisson law but with a random sign. S only contains binary relationships but Correct was only constrained to $\max(\hat{p}_1^j) = 5$. We used RMIXMOD to estimate the densities of each covariate. For each configuration, the walk was launched on 20 initial structures with a maximum of 9 000 steps each time.

4.2 Finding the structure

4.2.1 How to evaluate found structure?

The first criterion is ψ which is minimised in the MCMC. But in our case, ψ is based on the likelihood whose value don't have any intrinsic meaning. To show how far the found structure is from the true one in terms of S we define some indicators to compare the true model S and the found one \hat{S} . Global indicators:

- TL (True left): the number of found dependent variables that really are dependent $TL = |I_2 \cap \hat{I}_2|$
- WL (Wrong left) : the number of found dependent variables that are not dependent $WL = |\hat{I}_2| TL$
- ML (Missing left): the number of really dependent variables not found $ML = |I_2| TL$
- Δp_2 : the gap between the number of sub-regression in both model: $\Delta p_2 = |I_2| |\hat{I}_2|$. The sign defines if \hat{S} is too complex or too simple
- $\Delta compl$: the difference in complexity between both model: $\Delta compl = \sum_{j \in p_2} p_1^j \sum_{j \in \hat{p}_2} \hat{p}_1^j$

4.2.2 Results on S

In table 1 we compare found structures in different contexts with both Uniform (U) and Hierarchical Uniform (HU) a priori law on P(S). We see that (HU) hypothesis gives sparser models even if $\max(\hat{p}_1^j) = 5$. p = 40 so the maximum value for \hat{p}_2 is 20 in (HU) case and we see that this max is not reached. The stronger penalty implied by (HU) really is efficient. The datasets used for (U) and (HU) are the same to keep the comparison meaningful. These datasets and \hat{S} are those used for tables 2 to 9

It is also notable that (HU) has a greater computational cost than (U). But next versions of the package may optimize a bit more the penalization step and reduce this gap. We also notice that the MCMC is faster when there are numerous correlations (rejecting more candidates).

n	p_2	ψ	Time Mixmod	Time MCMC	TL	WL	ML	Δp_2	$\Delta compl$
30	0	U	0.4104	3.2428	0	5.43	0	-5.43	22.55
			(0.0275)	(0.3711)	(0)	(1.9346)	(0)	(1.9346)	(8.0884)
		HU	0.4104	8.2338	0	0.53	0	-0.53	2.27
			(0.0275)	(0.9045)	(0)	(0.7844)	(0)	(0.7844)	(3.3024)
30	16	U	0.4182	2.7735	10.96	5.93	4.98	-0.95	38.16
			(0.0329)	(0.1498)	(1.9844)	(2.0313)	(1.9948)	(0.9987)	(6.499)
		HU	0.4182	4.1876	11.61	4.57	4.33	-0.24	16.48
			(0.0329)	(0.2178)	(1.8743)	(1.9502)	(1.8752)	(0.4948)	(5.4892)
30	32	U	0.4456	2.9154	25.23	1.92	6.5	4.58	28
			(0.0429)	(0.1331)	(1.4761)	(1.0888)	(1.4668)	(0.9866)	(5.0831)
		HU	0.4456	4.0233	16.96	3.04	14.77	11.73	4.35
			(0.0429)	(0.1091)	(1.3993)	(1.3993)	(1.4692)	(0.5478)	(5.8833)
50	0	U	0.5229	4.7068	0	4.2	0	-4.2	13.35
			(0.0519)	(0.5865)	(0)	(1.7233)	(0)	(1.7233)	(5.6468)
		HU	0.5229	10.1198	0	0.13	0	-0.13	0.32
			(0.0519)	(0.5541)	(0)	(0.3667)	(0)	(0.3667)	(0.9732)
50	16	U	0.5205	3.3681	11.15	5.42	4.72	-0.7	22.85
			(0.0451)	(0.3123)	(1.93)	(1.9132)	(1.886)	(0.7317)	(5.7742)
		HU	0.5205	4.909	11.42	4.59	4.45	-0.14	7.55
			(0.0451)	(0.4556)	(1.9079)	(1.8968)	(1.8333)	(0.3487)	(4.0611)
50	32	U	0.5833	3.2683	28.17	1.38	3.7	2.32	12.74
			(0.0628)	(0.316)	(1.3711)	(0.9077)	(1.3143)	(0.8394)	(4.3359)
		HU	0.5833	4.3599	17.27	2.73	14.6	11.87	-2.61
			(0.0628)	(0.3729)	(1.1708)	(1.1708)	(1.2792)	(0.338)	(4.4854)
100	0	U	0.9623	12.9373	0	2.83	0	-2.83	6.23
			(0.077)	(1.7778)	(0)	(1.2953)	(0)	(1.2953)	(3.1999)
		HU	0.9623	20.9817	0	0.01	0	-0.01	0.02
			(0.077)	(1.9421)	(0)	(0.1)	(0)	(0.1)	(0.2)
100	16	U	1.1223	6.9647	11.67	4.8	4.25	-0.55	12.58
			(0.1122)	(0.5473)	(2.0003)	(2.0646)	(1.956)	(0.7833)	(3.9471)
		HU	1.1223	8.8486	12.04	3.95	3.88	-0.07	3.75
			(0.1122)	(0.7174)	(1.9223)	(1.9404)	(1.9137)	(0.2564)	(2.2625)
100	32	U	1.4343	5.9626	30.14	0.84	1.61	0.77	6.96
			(0.2528)	(0.3136)	(1.3928)	(0.8495)	(1.2941)	(0.7086)	(3.0975)
		HU	1.4343	7.3741	17.49	2.51	14.26	11.75	-3.76
			(0.2528)	(0.2748)	(1.1849)	(1.1849)	(1.2441)	(0.4794)	(4.4859)

Table 1: Results of the Markov chain with constraint $\hat{p}_1 \leq 5$. Mean observed and standard deviation (sd).

4.3 Results on prediction

4.3.1 Y depends on all variables in X

We first try the method with a response depending on all covariates (CORREG reduces the dimension and can't give the true model if there is a structure). The datasets used here were those from 1.

We observe that CORREG is better than classical methods especially whith the Hierarchical Uniform law. When the complexity of the true model is higher than $\frac{p}{2}$ Uniform hypothesis logically is better but (HU) still beats classical methods. We also observe in table 5 that simple methods like stepwise (here from the package LARS) can give good results in prediction.

n	p_2	ψ	indicator	OLS	CorReg \hat{S}	Correg S
30	0	U	MSE (sd)	262627.57 (732019)	5928332.4 (49005690.2)	262627.57 (732019)
			cpl (sd)	30 (0)	29.99 (0.1)	30 (0)
		HU	MSE (sd)	262627.57 (732019)	10381962.64 (90962496.9)	262627.57 (732019)
			cpl (sd)	30 (0)	30 (0)	30 (0)
30	16	U	MSE (sd)	510747.53 (2287539.8)	635.42 (335.2)	610.32 (424.9)
			cpl (sd)	30 (0)	24.11 (1)	25 (0)
		HU	MSE (sd)	510747.53 (2287539.8)	603.24 (415.6)	610.32 (424.9)
			cpl (sd)	30 (0)	24.82 (0.6)	25 (0)
30	32	U	MSE (sd)	178323.95 (1426610.2)	180.02 (44.9)	141.03 (27.8)
			cpl (sd)	30 (0)	13.85 (0.9)	9 (0)
		HU	MSE (sd)	178323.95 (1426610.2)	330.31 (134.8)	141.03 (27.8)
			cpl (sd)	30 (0)	21 (0)	9 (0)
50	0	U	MSE (sd)	528.08 (228.6)	886.54 (364.3)	528.08 (228.6)
			cpl (sd)	41 (0)	36.8 (1.7)	41 (0)
		HU	MSE (sd)	528.08 (228.6)	542.29 (235)	528.08 (228.6)
			cpl (sd)	41 (0)	40.87 (0.4)	41 (0)
50	16	U	MSE (sd)	612.72 (291.5)	239.17 (89.4)	200.32 (42.6)
			cpl (sd)	41 (0)	24.43 (0.7)	25 (0)
		HU	MSE (sd)	$612.72\ (291.5)$	207.6 (51.3)	200.32 (42.6)
			cpl (sd)	41 (0)	24.99(0.5)	25 (0)
50	32	U	MSE (sd)	$555.44\ (262.5)$	128.98 (18.1)	121.08 (11.9)
			cpl (sd)	41 (0)	11.45 (0.9)	9 (0)
		HU	MSE (sd)	$555.44 \ (262.5)$	171.9 (31.1)	121.08 (11.9)
			cpl (sd)	41 (0)	21 (0)	9 (0)
100	0	U	MSE (sd)	167.71 (20.6)	323.44 (124.1)	167.71 (20.6)
			cpl (sd)	41 (0)	38.17 (1.3)	41 (0)
		HU	MSE (sd)	167.71 (20.6)	167.98 (20.8)	167.71 (20.6)
			cpl (sd)	41 (0)	40.99 (0.1)	41 (0)
100	16	U	MSE (sd)	168.68 (22.4)	158.51 (51.6)	133.49 (12.2)
			cpl (sd)	41 (0)	24.53 (0.7)	25 (0)
		HU	MSE (sd)	168.68 (22.4)	137.03 (20.3)	133.49 (12.2)
			cpl (sd)	41 (0)	25.01 (0.4)	25 (0)
100	32	U	MSE (sd)	173.25 (22.9)	112.8 (10.7)	110.63 (6.9)
			cpl (sd)	41 (0)	10.02 (0.8)	9 (0)
		HU	MSE (sd)	173.25 (22.9)	127.4 (11.9)	110.63 (6.9)
			cpl (sd)	41 (0)	21 (0)	9 (0)

Table 2: OLS and OLS combined with constrained CorReg.Y depends on all variables in X. CorReg logically wins. When n < p the dataset was reduced to p = n automatically by a QR decomposition as the lm function of R does. Without selection, all models have $\min(n, p)$ non-zero coefficients.

n	p_2	ψ	indicator	LASSO	CorReg \hat{S}	Correg S
30	0	U	MSE (sd)	1246.35 (350.5)	1433.98 (526.7)	1246.35 (350.5)
			cpl (sd)	17.84 (5.5)	15.79 (5.6)	17.84 (5.5)
		HU	MSE (sd)	1246.35 (350.5)	1248.84 (341.4)	1246.35 (350.5)
			cpl (sd)	17.84 (5.5)	17.63 (5.6)	17.84 (5.5)
30	16	U	MSE (sd)	712.79 (405.2)	566.32 (226.4)	554.01 (257.1)
			cpl (sd)	16.68 (4.4)	15.35 (4.3)	16.14 (4.5)
		HU	MSE (sd)	712.79 (405.2)	551.92 (242.4)	554.01 (257.1)
			cpl (sd)	16.68 (4.4)	15.96 (4.4)	16.14 (4.5)
30	32	U	MSE (sd)	216.12 (128.3)	157.78 (39.4)	147.04 (28.4)
			$\operatorname{cpl}\left(\operatorname{sd}\right)$	11.23 (4.4)	8.56 (1.8)	7.91 (1.3)
		HU	MSE (sd)	216.12 (128.3)	178.88 (68.9)	147.04 (28.4)
			$\operatorname{cpl}\left(\operatorname{sd}\right)$	11.23 (4.4)	9.86(3)	7.91 (1.3)
50	0	U	MSE (sd)	658.38 (221.4)	872.32 (239.4)	658.38 (221.4)
			cpl (sd)	28.2(6)	22.95 (5.8)	28.2 (6)
		HU	MSE (sd)	658.38 (221.4)	652.72 (211.9)	658.38 (221.4)
			cpl (sd)	28.2 (6)	28.36(5.9)	28.2 (6)
50	16	U	MSE (sd)	274.34 (101.7)	268.76 (106.3)	225.64 (67.5)
			cpl (sd)	22.14(4.1)	19.52(2.9)	20.47(2.8)
		HU	MSE (sd)	274.34 (101.7)	233.8 (78.9)	225.64 (67.5)
			$\operatorname{cpl}\left(\operatorname{sd}\right)$	22.14 (4.1)	20.19(2.8)	20.47(2.8)
50	32	U	MSE (sd)	165.6 (73.7)	128.34 (18.8)	123.73 (12.8)
			$\operatorname{cpl}\left(\operatorname{sd}\right)$	12.93 (6.4)	8.51 (1.5)	8.09 (1.1)
		HU	MSE (sd)	165.6 (73.7)	135.84 (24.1)	123.73 (12.8)
			$\operatorname{cpl}\left(\operatorname{sd}\right)$	12.93 (6.4)	10.17 (3.1)	8.09 (1.1)
100	0	U	MSE (sd)	183.33 (31.1)	357.15 (133.4)	183.33 (31.1)
			cpl (sd)	37.78(2.4)	32.93(3.7)	37.78 (2.4)
		HU	MSE (sd)	183.33 (31.1)	183.78 (31.6)	183.33 (31.1)
			$\operatorname{cpl}\left(\operatorname{sd}\right)$	37.78(2.4)	37.75(2.4)	37.78 (2.4)
100	16	U	MSE (sd)	148.83 (18.6)	164.44 (54.3)	139.57 (16.1)
			$\operatorname{cpl}\left(\operatorname{sd}\right)$	25.22(4)	21.72(2)	22.3 (1.7)
		HU	MSE (sd)	148.83 (18.6)	142.97 (22.1)	139.57 (16.1)
			cpl (sd)	25.22 (4)	22.24 (1.8)	22.3 (1.7)
100	32	U	MSE (sd)	124.29 (19.9)	113.36 (11)	111.54 (7.2)
			$\operatorname{cpl}\left(\operatorname{sd}\right)$	13.18 (5.9)	8.68 (1.2)	8.53 (1)
		HU	MSE (sd)	124.29 (19.9)	118.33 (12)	111.54 (7.2)
			cpl (sd)	13.18 (5.9)	11.02 (2.9)	8.53 (1)

Table 3: LASSO (with LAR) combined with constrained CorReg.Y depends on all variables in X. CorReg logically wins. LASSO is better than OLS but is improved by CorReg even for large values of n.

n	p_2	ψ	indicator	Elasticnet	CorReg \hat{S}	Correg S
30	0	U	MSE (sd)	1 326.54 (388.4)	1 356.98 (331.4)	1 326.54 (388.4)
			cpl (sd)	12.14 (5.1)	12.5 (5.2)	12.14 (5.1)
		HU	MSE (sd)	1326.54 (388.4)	1307.96 (356.6)	1326.54 (388.4)
			cpl (sd)	12.14(5.1)	12.27(5.1)	12.14(5.1)
30	16	U	MSE (sd)	1 400.56 (1598.2)	668.57 (274.6)	653.59 (283.7)
			cpl (sd)	13.86 (6.8)	14.31 (5.7)	14.83 (5.3)
		HU	MSE (sd)	1 400.56 (1598.2)	643.25 (277.9)	653.59 (283.7)
			cpl (sd)	13.86 (6.8)	14.83 (5.6)	14.83 (5.3)
30	32	U	MSE (sd)	855.74 (582.6)	181.08 (51.9)	146.79 (32.7)
			cpl (sd)	15.57 (6.3)	11.19(2.4)	8.37 (1.3)
		HU	MSE (sd)	855.74 (582.6)	311.57 (163.9)	146.79 (32.7)
			cpl (sd)	15.57 (6.3)	14.87 (3.9)	8.37 (1.3)
50	0	U	MSE (sd)	738.94 (254.5)	914.56 (283.7)	738.94 (254.5)
			cpl (sd)	25.85 (8.7)	20.97(7.8)	25.85 (8.7)
		HU	MSE (sd)	738.94 (254.5)	751.06 (262.2)	738.94 (254.5)
			cpl (sd)	25.85 (8.7)	25.43 (8.8)	25.85 (8.7)
50	16	U	MSE (sd)	516.36 (226.6)	276.6 (133.6)	228.85 (89.3)
			cpl (sd)	26.72(7.4)	20.49(3.8)	21.8 (3.3)
		HU	MSE (sd)	516.36 (226.6)	239.77 (103.7)	228.85 (89.3)
			cpl (sd)	26.72 (7.4)	21.57(3.4)	21.8 (3.3)
50	32	U	MSE (sd)	328.7 (146.3)	130.61 (20.6)	124.01 (15.4)
			cpl (sd)	24.7 (7.1)	9.95 (1.5)	8.22 (1.2)
		HU	MSE (sd)	328.7 (146.3)	169.13 (40.2)	124.01 (15.4)
			cpl (sd)	24.7 (7.1)	16.04 (3.8)	8.22 (1.2)
100	0	U	MSE (sd)	175.93 (25.9)	355.79 (143.8)	175.93 (25.9)
			cpl (sd)	39.71 (1.7)	34.11 (4.6)	39.71 (1.7)
		HU	MSE (sd)	$175.93\ (25.9)$	$176.2\ (26.3)$	175.93 (25.9)
			cpl (sd)	39.71 (1.7)	39.69(1.7)	39.71 (1.7)
100	16	U	MSE (sd)	173.65 (25.1)	164.96 (59)	139.51 (17.6)
			cpl (sd)	35.54(4.2)	22.58(2.2)	23.26 (1.9)
		HU	MSE (sd)	173.65 (25.1)	143.04 (23.2)	139.51 (17.6)
			cpl (sd)	35.54(4.2)	23.16 (1.9)	23.26 (1.9)
100	32	U	MSE (sd)	161.92 (25.2)	113.61 (10.9)	111.46 (7.2)
			cpl (sd)	30.78 (6.4)	9.25(1.2)	8.67 (1)
		HU	MSE (sd)	161.92 (25.2)	127.44 (13.3)	111.46 (7.2)
			cpl (sd)	30.78 (6.4)	17.85 (2.7)	8.67 (1)

Table 4: Elastic net combined with constrained $\mathrm{CorReg}.Y$ depends on all variables in X. CorReg logically wins. LASSO was better.

\overline{n}	p_2	ψ	indicator	Stepwise	CorReg \hat{S}	$\operatorname{CorReg} S$
30	0	U	MSE (sd)	1 919.43 (861.8)	2 014.13 (683.6)	1 919.43 (861.8)
			cpl (sd)	22.8 (3.3)	19.34 (4.9)	22.8 (3.3)
		HU	MSE (sd)	1 919.43 (861.8)	1 885.39 (814)	1 919.43 (861.8)
			cpl (sd)	22.8 (3.3)	22.73 (3.1)	22.8 (3.3)
30	16	U	MSE (sd)	696.46 (492.9)	662.07 (285.9)	660.93 (354)
			cpl (sd)	15.9(3.8)	15.1 (3.7)	15.76(3.7)
		HU	MSE (sd)	696.46 (492.9)	655.94 (325.7)	660.93 (354)
			cpl (sd)	15.9(3.8)	15.72(3.6)	15.76 (3.7)
30	32	U	MSE (sd)	394.93 (356)	155.18 (36.4)	147.81 (30.3)
			cpl (sd)	16.1 (6.3)	8.26 (1.7)	7.77 (1.3)
		HU	MSE (sd)	394.93 (356)	189.49 (79)	147.81 (30.3)
			cpl (sd)	16.1 (6.3)	10.23 (3)	7.77 (1.3)
50	0	U	MSE (sd)	745.88 (241.7)	962.12 (319.3)	745.88 (241.7)
			cpl (sd)	27.19(4.6)	22.9(4.5)	27.19 (4.6)
		HU	MSE (sd)	745.88 (241.7)	749.08 (244.2)	745.88 (241.7)
			cpl (sd)	27.19(4.6)	27.21(4.5)	27.19 (4.6)
50	16	U	MSE (sd)	279.82 (151.1)	290.13 (117.3)	239.61 (69.5)
			cpl (sd)	21.24 (5.2)	18.13 (2.8)	19.23 (2.7)
		HU	MSE (sd)	279.82 (151.1)	252.35 (108.5)	239.61 (69.5)
			cpl (sd)	21.24 (5.2)	18.97 (2.7)	19.23 (2.7)
50	32	U	MSE (sd)	199.3 (158.7)	126.81 (17.4)	124.19 (13)
			cpl (sd)	14.31 (6.2)	8.16 (1.2)	8.04 (1)
		HU	MSE (sd)	199.3 (158.7)	136.32 (25.2)	124.19 (13)
			cpl (sd)	14.31 (6.2)	9.59 (2.6)	8.04 (1)
100	0	U	MSE (sd)	193.26 (34.9)	387.18 (151.6)	193.26 (34.9)
			cpl (sd)	36.3(2.5)	30.94 (4.2)	36.3(2.5)
		HU	MSE (sd)	193.26 (34.9)	194.43 (37.3)	193.26 (34.9)
			cpl (sd)	36.3(2.5)	36.22 (2.7)	36.3 (2.5)
100	16	U	MSE (sd)	145.46 (20.1)	165.74 (56.1)	140.27 (17.5)
			cpl (sd)	23.44(3.8)	21.23(2.1)	21.93(1.7)
		HU	MSE (sd)	145.46 (20.1)	143.74 (23.2)	140.27 (17.5)
			cpl (sd)	23.44 (3.8)	21.86 (1.7)	21.93 (1.7)
100	32	U	MSE (sd)	128.44 (18.4)	113.27 (10.7)	111.77 (7.2)
			cpl (sd)	13.58 (5.2)	8.59 (1.1)	8.48 (1)
		HU	MSE (sd)	128.44 (18.4)	118.61 (11.5)	111.77 (7.2)
			cpl (sd)	13.58 (5.2)	10.75 (2.8)	8.48 (1)

Table 5: Stepwise combined with constrained CorReg. Y depends on all variables in X. CorReg logically wins but stepwise is quite good compared to elasticnet.

4.3.2 Y depends only on covariates in X_2 (worst case for us)

We now try the method with a response depending only on variables in X_2 . The datasets used here were still those from 1. Depending only on X_2 imply sparsity and impossibility to obtain the true model when using the true structure. Correct is still better than classical methods.

n	p_2	ψ	indicator	OLS	CorReg \hat{S}	CorReg S
30	16	Ü	MSE (sd)	81781.54 (247281.7)	507.1 (269.2)	593.03 (428.3)
			cpl (sd)	30 (0)	24.11 (1)	25 (0)
		HU	MSE (sd)	81781.54 (247281.7)	587.07 (425.6)	593.03 (428.3)
			cpl (sd)	30 (0)	24.82 (0.6)	25 (0)
30	32	U	MSE (sd)	74136.81 (297716.1)	189.45 (48.1)	144.51 (31.4)
			cpl (sd)	30 (0)	13.85 (0.9)	9 (0)
		HU	MSE (sd)	74136.81 (297716.1)	340.53 (169.4)	144.51 (31.4)
			cpl (sd)	30 (0)	21 (0)	9 (0)
50	16	U	MSE (sd)	684.04 (438.1)	190.97 (37.7)	197.32 (40)
			cpl (sd)	41 (0)	24.43 (0.7)	25 (0)
		HU	MSE (sd)	684.04 (438.1)	196.36 (40.5)	197.32 (40)
			cpl (sd)	41 (0)	24.99(0.5)	25 (0)
50	32	U	MSE (sd)	596.32 (323)	126.94 (15.5)	119.39 (12.4)
			cpl (sd)	41 (0)	11.45 (0.9)	9 (0)
		HU	MSE (sd)	596.32(323)	168.03 (27.7)	119.39 (12.4)
			cpl (sd)	41 (0)	21 (0)	9 (0)
100	16	U	MSE (sd)	168.35 (20.1)	133.75 (12.4)	135.08 (13.1)
			cpl (sd)	41 (0)	24.53 (0.7)	25 (0)
		HU	MSE (sd)	168.35 (20.1)	135.04 (13.1)	135.08 (13.1)
			cpl (sd)	41 (0)	25.01 (0.4)	25 (0)
100	32	U	MSE (sd)	168.07 (21.4)	109.61 (7.4)	108.64 (7.2)
			cpl (sd)	41 (0)	10.02 (0.8)	9 (0)
		HU	MSE (sd)	168.07 (21.4)	124.23 (11.1)	108.64 (7.2)
			cpl (sd)	41 (0)	21 (0)	9 (0)

Table 6: OLS and OLS combined with constrained CORREG.Y depends on all variables in X_2 . Sometimes \hat{S} gives better results than S because S is penalized by the fact that it relies only on covariates not in the true model.

n	p_2	ψ	indicator	LAR	CorReg \hat{S}	Correg S
30	16	U	MSE (sd)	367.45 (198.4)	298.32 (122.4)	292.53 (102.2)
			cpl (sd)	14.37(5.1)	12.9(3.8)	12.84(3.8)
		HU	MSE (sd)	367.45 (198.4)	298.1 (119.1)	292.53 (102.2)
			cpl (sd)	14.37 (5.1)	12.81 (3.8)	12.84(3.8)
30	32	U	MSE (sd)	273.12 (267.2)	166.93 (49)	152.45 (39.5)
			cpl (sd)	11.7 (5.8)	8.7(2.5)	7.45(1.4)
		HU	MSE (sd)	273.12 (267.2)	175.68 (57.5)	152.45 (39.5)
			cpl (sd)	11.7 (5.8)	9.19 (2.9)	7.45(1.4)
50	16	U	MSE (sd)	189.64 (52.5)	171.97 (45.9)	176.58 (46.8)
			cpl (sd)	14.15 (3.9)	13.4(2.8)	13.5(3.1)
		HU	MSE (sd)	189.64 (52.5)	174.36 (45.3)	$176.58 \ (46.8)$
			cpl (sd)	14.15 (3.9)	13.57(3)	13.5(3.1)
50	32	U	MSE (sd)	163.88 (67.4)	124.53 (17)	121.99 (16.6)
			cpl (sd)	13.06 (7.2)	8.29 (1.6)	7.73(1.2)
		HU	MSE (sd)	163.88 (67.4)	133.7 (26.4)	121.99 (16.6)
			cpl (sd)	13.06 (7.2)	9.65(3.3)	7.73(1.2)
100	16	U	MSE (sd)	129.65 (14.8)	127.11 (12.1)	127.46 (12.2)
			cpl (sd)	14.86 (3.1)	13.97(2.5)	14.04 (2.5)
		HU	MSE (sd)	129.65 (14.8)	127.46 (12.2)	127.46 (12.2)
			cpl (sd)	14.86 (3.1)	14.02 (2.5)	14.04 (2.5)
100	32	U	MSE (sd)	121.21 (20.1)	109.66 (7.9)	109.04 (7.8)
			cpl (sd)	12.84 (5.2)	8.42 (1.1)	8.11 (0.8)
		HU	MSE (sd)	121.21 (20.1)	112.56 (11.2)	109.04 (7.8)
			cpl (sd)	12.84 (5.2)	9.77(2.3)	8.11 (0.8)

Table 7: LASSO (with LAR) combined with constrained CorReg.Y depends on all variables in X_2 .

n	p_2	ψ	indicator	Elasticnet	CorReg \hat{S}	Correg S
30	16	U	MSE (sd)	499.32 (218.8)	311.41 (137.4)	305.09 (123.5)
			cpl (sd)	14.08 (6.1)	11.75 (4.8)	11.49 (4.7)
		HU	MSE (sd)	499.32 (218.8)	307.6 (128.7)	305.09 (123.5)
			cpl (sd)	14.08 (6.1)	11.63 (4.7)	11.49 (4.7)
30	32	U	MSE (sd)	691.55 (503.2)	193.45 (54.2)	150.17 (35)
			cpl (sd)	12.54 (7.2)	10.26 (2.7)	7.81 (1.4)
		HU	MSE (sd)	691.55 (503.2)	279.21 (117.3)	150.17 (35)
			cpl (sd)	12.54 (7.2)	13 (4.3)	7.81 (1.4)
50	16	U	MSE (sd)	282.05 (134.8)	180.64 (47.6)	181.49 (48.6)
			cpl (sd)	20.37(6.3)	13.68 (4)	14.05 (4)
		HU	MSE (sd)	282.05 (134.8)	180.84 (47.5)	181.49 (48.6)
			cpl (sd)	20.37(6.3)	14.03(4)	14.05 (4)
50	32	U	MSE (sd)	308.13 (142.9)	129.39 (19.6)	122.71 (17.8)
			cpl (sd)	22.94(7.7)	9.37(2.1)	7.91 (1.4)
		HU	MSE (sd)	308.13 (142.9)	159.14 (30.4)	122.71 (17.8)
			cpl (sd)	$22.94\ (7.7)$	15.42(4)	7.91 (1.4)
100	16	U	MSE (sd)	150.14 (19.7)	127.48 (14)	128.21 (15)
			cpl (sd)	25.6(5.4)	14.73(3.8)	14.73 (3.9)
		HU	MSE (sd)	150.14 (19.7)	128.21 (14.9)	128.21 (15)
			cpl (sd)	25.6(5.4)	14.7 (3.9)	14.73 (3.9)
100	32	U	MSE (sd)	157.18 (23.5)	110.24 (8)	109.24 (7.8)
			cpl (sd)	30.41 (6.7)	8.96 (1.3)	8.3 (1)
		HU	MSE (sd)	$157.18\ (23.5)$	123.8 (12.5)	109.24 (7.8)
			$\operatorname{cpl}\left(\operatorname{sd}\right)$	30.41 (6.7)	17.18 (3.3)	8.3 (1)

Table 8: Elastic net (with Elasticnet) combined with constrained CorReg. Y depends on all variables in X_2 .

n	p_2	ψ	indicator	Stepwise	CorReg \hat{S}	Correg S
30	16	U	MSE (sd)	394.59 (454.6)	341.47 (153.2)	351.14 (148)
			cpl (sd)	13.09 (3.2)	12.9(2.9)	12.92 (3)
		HU	MSE (sd)	394.59 (454.6)	353.99 (157)	351.14 (148)
			cpl (sd)	13.09(3.2)	12.83(2.9)	12.92 (3)
30	32	U	MSE (sd)	410.11 (343.5)	163.05 (45.4)	152.08 (38.1)
			cpl (sd)	15.8 (6.5)	8.29(2.2)	7.34 (1.3)
		HU	MSE (sd)	410.11 (343.5)	178.94 (69.7)	152.08 (38.1)
			cpl (sd)	15.8 (6.5)	8.89(3)	7.34 (1.3)
50	16	U	MSE (sd)	180.67 (52.7)	173.02 (38.3)	174.84 (36.8)
			cpl (sd)	13.56 (3.7)	12.78(2.4)	13.07(2.4)
		HU	MSE (sd)	180.67 (52.7)	175.3 (38.1)	174.84 (36.8)
			cpl (sd)	13.56 (3.7)	13.01 (2.4)	13.07(2.4)
50	32	U	MSE (sd)	188.91 (88.5)	124.38 (17.4)	122.38 (16.7)
			cpl (sd)	14.74 (6.6)	8.07(1.6)	7.65(1.2)
		HU	MSE (sd)	188.91 (88.5)	134.22 (23.6)	122.38 (16.7)
			cpl (sd)	14.74 (6.6)	9.5(2.8)	7.65 (1.2)
100	16	U	MSE (sd)	128.18 (13.7)	127.43 (13.7)	127.71 (13.3)
			cpl (sd)	13.78(2.8)	13.51(2)	13.59(2.1)
		HU	MSE (sd)	128.18 (13.7)	$127.71 \ (13.3)$	127.71 (13.3)
			cpl (sd)	13.78(2.8)	13.59(2.1)	13.59(2.1)
100	32	U	MSE (sd)	126.15 (20.6)	109.47(8)	109.04 (7.8)
			cpl (sd)	13.25 (4.4)	8.3(1)	8.12 (0.8)
		HU	MSE (sd)	126.15 (20.6)	113.96 (11.4)	109.04 (7.8)
			cpl (sd)	13.25 (4.4)	10.08 (2.6)	8.12 (0.8)

Table 9: Stepwise combined with constrained CorReg. Y depends on all variables in X_2 .

4.4 Robustess of the model

We have generated non linear structures (Tables 10 and 11). Variables in X_2 depends on the log or the square (randomly choosen with equiprobability) of a variable in X_1 . Dependencies are still real but non linear. Correct still found dependencies. One example of non-linear structure found with Correct on real datasets is given in section 5.1.

n	p_2	ψ	Time Mixmod	Time MCMC	TL	WL	ML	Δp_2	$\Delta compl$
30	16	U	0.4313	2.7769	9.96	2.45	5.95	3.5	29.42
			(0.0391)	(0.0883)	(1.3175)	(1.3808)	(1.3056)	(1.453)	(6.2639)
		HU	0.4313	4.9079	7.28	1.25	8.63	7.38	5.77
			(0.0391)	(0.4353)	(1.5769)	(0.9987)	(1.5548)	(1.6316)	(6.1297)

Table 10: Results of the Markov chain for non linear structure (log and square). Mean observed and standard deviation (sd).

n	p_2	ψ	indicator	LASSO	CorReg \hat{S}	CorReg S
30	0	U	MSE (sd)	996.06 (323.7)	1100.17 (391.4)	1501.25 (905.9)
			cpl (sd)	17.66 (5.4)	14.7 (4.7)	11.97(4.9)
		HU	MSE (sd)	996.06 (323.7)	1094.5 (621.8)	1501.25 (905.9)
			cpl (sd)	17.66 (5.4)	16.98 (4.8)	11.97 (4.9)

Table 11: Y depends on all variables in X. CorReg is not too far from LASSO even if it stays behind.

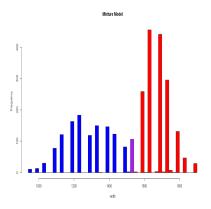
5 Numerical results on real datasets

5.1 Quality case study

This work takes place in steel industry context, with quality oriented objective: to understand and prevent quality problems on finished product, knowing the whole process.

We have :

- a quality parameter (confidential) as response variable,
- 205 variables from the whole process to explain it.
- The stakes: a hundred euros per ton (for information: Dunkerque's site aims to produce up to 7.5 millions tons a year)



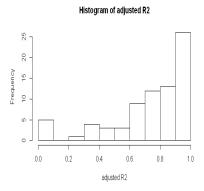




Figure 4: Example of non-Gaussian real variable easily modeled by a Gaussian mixture

Figure 5: R_{adj}^2 of the 76 sub-regressions.

Figure 6: Histogram of correlations in X.

We get a training set of n=3000 products described by p=205 variables from the industrial process and a validation sample of 847 products. Let's note ρ the absolute value of correlations between two covariates. Industrial variables are naturally highly correlated as the width and the weight of a steel slab ($\rho=0.905$), the temperature before and after some tool ($\rho=0.983$), the roughness of both faces of the product ($\rho=0.919$), a mean and a max ($\rho=0.911$). Correct also found more complex structures describing physical models, like Width = f (Mean.flow , Mean.speed.CC) even if the true Physcial model is not linear : Width = flow / (speed * thickness) (here thickness is constant). Non linear regulation models used to optimize the process were also found (but are confidential). These first results are easily understandable and meet metallurgists expertise. The algorithm gives a structure of $p_2=76$ subregressions with a mean of $p_1=5.17$ regressors. In $p_2=76$ subregressions with a mean of $p_3=5.17$ regressors. In $p_4=5.17$ the number of $p_4=5.17$ smaller than in $p_4=5.17$ regressors.

It is now time to look at the predictive results (Figure 5.1). The best model found when not using Correct is given by the LASSO. But when using Correct elasticnet produces a better model in terms of prediction. LASSO gives a model with 21 non-zero coefficients and elasticnet with Correct gives a model with 40 non-zero parameters but 6.40% better in prediction on the validation sample (847 products). 14 non-zero coefficients are common between the two models. Elasticnet alone get a model with 78 parameters that is improved by 9.75% in prediction when used with Correct When using LASSO with Correct we obtain a model with 24 non-zero coefficients that is 4.11% better than LASSO alone. We also computed the OLS model (without selection) and the naive one (estimating the response by the mean of the learning set). All the MSE were modified here to obtain a value of 100 for the best (to preserve confidentiality). Elasticnet with Correct is 13.51% better than OLS.

In terms of interpretation, the main regression comes with the family of regression so it gives a better understanding of the consequences of corrective actions on the whole process. It typically

Predictive error (MSE) on a validation sample

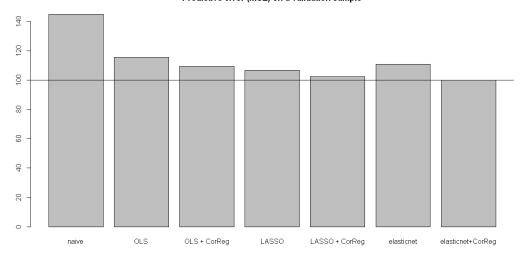


Figure 7: MSE comparison on industrial dataset. Learning set : 3~000 products, validation set : 847 products

Model	MSE	Complexity (with intercept)
OLS	115.63	206
Correg + OLS	109.59	130
LASSO	106.84	21
CorReg + LASSO	102.45	24
elasticnet	110.81	78
Corred + elasticnet	100	40

Table 12: Results obtained on a validation sample.

permits to determine the *tuning parameters* whereas LASSO would point variables we can't directly act on. So it becomes easier to take corrective actions on the process to reach the goal. The stakes are so important that even a little improvement leads to consequent benefits, and we don't even talk of the impact on the market shares that is even more important.

5.2 Production case study

This second example is about a phenomenon that impacts the productivity of a steel plan. We have:

- a (confidential) response variable,
- p = 145 variables from the whole process to explain it but only n = 100 individuals.
- The stakes : 20% of productivity to gain

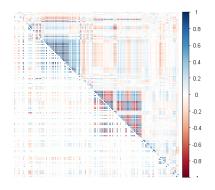


Figure 8: Correlations between the covariates in X (upper) and \hat{X}_1 (lower).

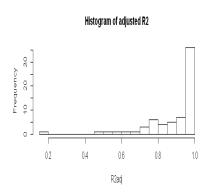


Figure 9: R_{adj}^2 of the 67 sub-regressions.



Figure 10: Histogram of correlations in X.

Here n < p so we only compare the leave-one-out cross-validation MSE. Corresponding improves LASSO by 5.24% and elasticated by 8.60%. Corresponding to the leave-one-out MSE. In this precise case, Corresponding to the leave-one-out MSE. In this precise case, Corresponding to the leave-one-out MSE.

Model	MSE	Complexity (with intercept)
LASSO	105.54	34
CorReg + LASSO	100	18
elasticnet	129.94	13
Corred + elasticnet	118.76	21

Table 13: Results obtained with leave-one out cross-validation. n = 100, p = 145.

covariates in interpretation and to find the relevant part of the process to optimize.

6 Conclusion and perspectives

We have seen that correlations can lead to serious estimation and variable selection problems in linear regression and that in such a context, it can be useful to explicitly model the structure between the covariates and to use this structure (even sequentially) to avoid correlations issues. We also show that real industrial context faces this kind of situations so our model can help to interprete and predict physical phenomenon efficiently and to help to manage missing values. But for now we still need a full dataset to learn the structure between the covariates and even if correlations are strong, some information is lost. Further work is needed to face these two challenges.

Correct is accessible on CRAN and has already proved its efficiency on real regression problematics in industry. Correct is strength is its great interpretability of the model, composed of several short linear regression easily managed by non-statisticians while strongly reducing correlations issues that are everywhere in industry. Nevertheless, we need to enlarge its application field to missing values, also very commons in industry. The actual generative model allows such a functionality without supplementary hypothesis and this also is a strength of Correct.

Another perspective would be to take back lost information (the residual of each sub-regression) to improve predictive efficiency when needed. It would only consists in a second step of linear regression between the residuals and would thus still be able to use any selection method.

This paper only treats linear regression but such a pretreatment could be used for logistic regression, *etc.* So the subject is still wide opened.

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7 Appendices

7.1 Identifiability of the structure

The model presented above relies on a discrete structure S between the covariates. But to find it we need identifiability property to insure the MCMC will asymptotically find the true model. Identifiability of the discrete structure is asked in following terms: Is it possible to find another structure \tilde{S} of linear regression between the covariates leading to the same joint distribution and marginal distributions? The answer is no. Thus S is identifiable.

7.1.1 Alternative neighbourhoods for the MCMC

We have here at each step $|\mathcal{V}_{S,j}| = p$ candidates but some other constraints can be added on the definition of S and will consequently modify the size of the neighbourhood (for example a maximum complexity for the internal regressions or the whole structure, a maximum number of internal regressions, etc.). Correct allows to modify this neighbourhood to better fit users constraints. Relaxation (column-wise and row-wise) is optional but gives more stability to the number of feasible candidates at each step and allows to modify several parts of I_1 in only one step when needed. Hence it improves efficiency by a significant reinforcement of the irreductibility of the Markov chain. Rejecting candidates instead of doing the relaxation steps will however reduce the number of evaluated candidates and thus accelerate the walk. So it can be used for a warming phase when n is great and time is missing.

The hierchical uniform hypothesis made above for P(S) implies $p_2 < \frac{p}{2}$ and $p_1^j < \frac{p}{2}$ so candidates may be rejected to satisfy this hypothesis. Stronger constraints on p_2 and/or p_1 can be given in CORREG if relevant.

If the algorithm did not have time to converge (stationnarity), it can be continued with a few step for which the neighboorhood would only contain smaller candidates (in terms of complexity). It is equivalent to ask for each element in I_1 if the criterion ψ would be better without it. Thus it can be seen as a final cleaning step. But in fact, it's just continuing the MCMC with a reduced neighbourhood.