

Supplementary material: fast structure learning with modular regularization

A Proofs

A.1 Proof of proposition 2.1

Proposition. 2.1 restated. The random variables X and Z are described by a directed graphical model where the parents of X are in Z and the Z 's are independent if and only if $TC(X|Z) + TC(Z) = 0$.

Proof. Because TC is always non-negative,

$$TC(X|Z) + TC(Z) = 0 \Leftrightarrow TC(Z) = 0 \text{ and } TC(X|Z) = 0.$$

We also have the following standard statements [6]

$$TC(X|Z) = 0 \Leftrightarrow \forall x, z, p(x|z) = \prod_{i=1}^p p(x_i|z),$$

$$TC(Z) = 0 \Leftrightarrow \forall z, p(z) = \prod_{j=1}^m p(z_j).$$

Putting these together, we have

$$\forall x, z, p(x, z) = \prod_{i=1}^p \prod_{j=1}^m p(x_i|z)p(z_j).$$

We can see that this statement is equivalent to the definition of a Bayesian network for random variables X, Z with respect to the graph in Fig. 1a. \square

A.2 Proof of theorem 2.1

Theorem. 2.1 restated. A multivariate Gaussian distribution $p(x, z)$ is a modular latent factor model if and only if $TC(X|Z) + TC(Z) = 0$ and $\forall i, TC(Z|X_i) = 0$.

Proof. First we show that for any modular latent factor model, even non-Gaussian, the constraints are satisfied. Thm. 2.1 establishes that the model implies $TC(X|Z) + TC(Z) = 0$. We must show that the additional restriction that each X_i has only one parent, Z_{π_i} , implies the condition $\forall i, TC(Z|X_i) = 0$. Looking at the rules for d-separation we see that Z_1, \dots, Z_m are independent conditioned on X_i . Therefore, $\forall i, TC(Z|X_i) = 0$.

Now, we show that a multivariate Gaussian distribution $p(x, z)$ with $TC(X|Z) + TC(Z) = 0$ and $\forall i, TC(Z|X_i) = 0$ is a modular latent factor model:

$$\forall x, z, p(x, z) = \prod_{i=1}^p p(x_i|z_{\pi_i}) \prod_{j=1}^m p(z_j), \text{ for some } \pi_i \in \{1, 2, \dots, m\}.$$

By Thm. 2.1 we have that $\forall x, z, p(x, z) = \prod_{i=1}^p p(x_i|z) \prod_{j=1}^m p(z_j)$. To complete the proof we show that $(TC(Z) = 0 \ \& \ TC(Z|X_i) = 0) \Rightarrow p(x_i|z) = p(x_i|z_{\pi_i})$ for some $\pi_i \in \{1, \dots, m\}$. We have

$$\begin{aligned} p(x_i|z) &= p(x_i)/p(z) \prod_{j=1}^m p(z_j|x_i) \\ &= p(x_i) \prod_{j=1}^m p(z_j|x_i)/p(z_j) \\ &= p(x_i) \prod_{j=1}^m p(z_j, x_i)/(p(x_i)p(z_j)). \end{aligned} \tag{5}$$

We also have that $TC(Z|X_i) = 0 \Rightarrow \forall j \neq k, \text{Cov}[Z_j, Z_k|X_i] = 0$. For Gaussians $\text{Cov}[Z_j, Z_k|X_i] = \text{Cov}[Z_j, Z_k] - \text{Cov}[Z_j, X_i]\text{Cov}[Z_k, X_i]/\text{Var}[X_i]$. Having $\text{Var}[X_i] > 0$ and ($TC(Z) = 0 \Rightarrow \text{Cov}[Z_j, Z_k] = 0$), we get $\text{Cov}[Z_j, X_i] = 0 \vee \text{Cov}[Z_k, X_i] = 0$. Therefore, for all but at most one index, π_i , it must be the covariance of X_i and Z_j is zero, so that $p(z_j, x_i) = p(x_i)p(z_j)$. Putting this in Eq. (5) we get $p(x_i|z) = p(x_i|z_{\pi_i})$.

Note that we cannot remove the Gaussian assumption, since it is possible to have $TC(X|Z) = 0, TC(Z) = 0$, and $\forall i, TC(Z|X_i) = 0$, but still have two non-trivial parents for one X_i . For example, if $Z_1, Z_2 \stackrel{iid}{\sim} \text{Bernoulli}(1/2)$ and $X_1 = 2Z_1 + Z_2$. It can be easily seen that the conditions are satisfied, but it is impossible to model X_1 with only Z_1 or Z_2 as its parent. \square

B Complete derivation of linear CorEx

In this section we describe the complete derivation of linear CorEx. The first step is to define the family of joint distributions we are searching over by parametrizing $p_W(z|x)$. If $X_{1:p}$ is Gaussian, then we can ensure $X_{1:p}, Z_{1:m}$ are jointly Gaussian by parametrizing $p_W(z_j|x) = \mathcal{N}(w_j^T x, \eta_j^2)$, $w_j \in \mathbb{R}^p$, $j = 1..m$, or equivalently by $z = Wx + \epsilon$ with $W \in \mathbb{R}^{m \times p}$, $\epsilon \sim \mathcal{N}(0, \text{diag}(\eta_1^2, \dots, \eta_m^2))$. The noise variances η_j^2 are taken to be constants. Please note the implicit conditional independence assumption, $TC(Z|X) = 0$, we are making using this parameterization. We do this assumption since modular latent factor models have $TC(Z|X) = 0$, and it simplifies further derivations. W.l.o.g. we assume the data is standardized so that $\mathbb{E}[X_i] = 0, \mathbb{E}[X_i^2] = 1$ ¹. If it is not standardized we can standardize it using the empirical means and standard deviations. Motivated by Thm. 2.1, we will start with the following optimization problem:

$$\underset{W}{\text{minimize}} TC(X|Z) + TC(Z) + \sum_{i=1}^p Q_i, \quad (6)$$

where Q_i are regularization terms for encouraging modular solutions (i.e. encouraging solutions with smaller value of $TC(Z|X_i)$)². We will later specify this regularizer as a non-negative quantity that goes to zero in the case of exactly modular latent factor models. The $TC(X|Z) + TC(Z)$ part of the Eq. (6) can be rewritten as follows:

$$\begin{aligned} TC(X|Z) + TC(Z) &= \sum_{i=1}^p H(X_i|Z) - H(X|Z) + \sum_{j=1}^m H(Z_j) - H(Z) \\ &= \sum_{i=1}^p H(X_i|Z) + \sum_{j=1}^m H(Z_j) - (H(X|Z) + H(Z)) \\ &= \sum_{i=1}^p H(X_i|Z) + \sum_{j=1}^m H(Z_j) - (H(Z|X) + H(X)) \\ &= \sum_{i=1}^p H(X_i|Z) + \sum_{j=1}^m (H(Z_j) - H(Z_j|X)) + H(X) \\ &\propto \sum_{i=1}^p H(X_i|Z) + \sum_{j=1}^m I(Z_j; X). \end{aligned} \quad (7)$$

The first two lines invoke definitions and re-arrange. The third line uses Bayes' rule to rewrite the entropies. The fourth line invokes conditional independence of Z 's conditioned on X . Next, we write

¹Unless specified all expectations are taken with respect to the joint distribution $p_W(x, z)$.

²One can set $Q_i \propto TC(Z|X_i)$. However, we choose not to do this since we do not have a derivation that leads the resulting objective into an equivalent, but efficiently computable objective.

out the explicit form of expressions in Eq. (7) for Gaussians and ignore constants:

$$\begin{aligned}
& \sum_{i=1}^p H(X_i|Z) + \sum_{j=1}^m I(Z_j; X) \\
&= \sum_{i=1}^p \frac{1}{2} \mathbb{E}_Z \log (2\pi e \text{Var}[X_i|Z]) + \sum_{j=1}^m (H(Z_j) - H(Z_j|X)) \\
&= \sum_{i=1}^p \frac{1}{2} \log \mathbb{E}_Z [2\pi e \text{Var}[X_i|Z]] + \sum_{j=1}^m (H(Z_j) - H(Z_j|X)) \\
&\propto \frac{1}{2} \sum_{i=1}^p \log \mathbb{E} [(X_i - \mathbb{E}_{X_i|Z}[X_i|Z])^2] + \frac{1}{2} \sum_{j=1}^m (\log \text{Var}[Z_j] - \mathbb{E}_X \log \text{Var}[Z_j|X]) \\
&\propto \frac{1}{2} \sum_{i=1}^p \log \mathbb{E} [(X_i - \mathbb{E}_{X_i|Z}[X_i|Z])^2] + \frac{1}{2} \sum_{j=1}^m (\log \mathbb{E} [Z_j^2] - \log(\eta_j^2)) \\
&\propto \frac{1}{2} \sum_{i=1}^p \log \mathbb{E} [(X_i - \mathbb{E}_{X_i|Z}[X_i|Z])^2] + \frac{1}{2} \sum_{j=1}^m \log \mathbb{E} [Z_j^2]. \tag{8}
\end{aligned}$$

We used the fact that the differential entropy of a Gaussian variable with variance σ^2 is equal to $1/2 \log(2\pi e \sigma^2)$. Also, we used the fact that if A, B are jointly Gaussian random variables, then $H(A|B) \propto \mathbb{E}_B \log \text{Var}[A|B] = \log \mathbb{E}_B \text{Var}[A|B]$. The logarithm and expectation can be swapped because for Gaussians $\text{Var}[A|B]$ is constant for any value of B . In the fifth line we replace $\text{Var}[Z_j]$ with $\mathbb{E} [Z_j^2]$, because having $\mathbb{E} [X] = 0$ and $z_j = w_j^T x + \epsilon_j$ implies $\mathbb{E} [Z_j] = 0$. Considering Eq. (8), the problem (6) becomes:

$$\underset{W}{\text{minimize}} \sum_{i=1}^p (1/2 \log \mathbb{E} [(X_i - \mu_{X_i|Z})^2] + Q_i) + \sum_{j=1}^m 1/2 \log \mathbb{E} [Z_j^2], \tag{9}$$

where $\mu_{X_i|Z} = \mathbb{E}_{X_i|Z}[X_i|Z]$. For Gaussians, calculating $\mu_{X_i|Z}$ requires a computationally undesirable matrix inversion. Instead, we will select Q_i to eliminate this term while also encouraging modular structure. According to Thm. 2.1 modular models obey $TC(Z|X_i) = 0$, which implies that $p(x_i|z) = p(x_i)/p(z) \prod_j p(z_j|x_i)$. Let $\nu_{X_i|Z}$ be the conditional mean of X_i given Z under such factorization. Then it will have the following form (see Sec. B.1 for the derivation):

$$\begin{aligned}
\nu_{X_i|Z} &= \frac{1}{1 + r_i} \sum_{j=1}^m \frac{Z_j B_{j,i}}{\sqrt{\mathbb{E} [Z_j^2]}}, \\
\text{with } R_{j,i} &= \frac{\mathbb{E} [X_i Z_j]}{\sqrt{\mathbb{E} [X_i^2] \mathbb{E} [Z_j^2]}}, B_{j,i} = \frac{R_{j,i}}{1 - R_{j,i}^2}, r_i = \sum_{j=1}^m R_{j,i} B_{j,i}.
\end{aligned}$$

We see that computing $\nu_{X_i|Z}$ is easier since it requires no matrix inversion and depends only on pairwise statistics between observed and latent variables. If we let $Q_i = 1/2 \log \mathbb{E} [(X_i - \nu_{X_i|Z})^2] - 1/2 \log \mathbb{E} [(X_i - \mu_{X_i|Z})^2]$, we will replace $\mu_{X_i|Z}$ with $\nu_{X_i|Z}$ in problem (9). To see why this also

encourages modular structures we note that

$$\begin{aligned}
Q_i &= \frac{1}{2} \log \mathbb{E} [(X_i - \nu_{X_i|Z})^2] - \frac{1}{2} \log \mathbb{E} [(X_i - \mu_{X_i|Z})^2] \\
&= \frac{1}{2} \log \frac{\mathbb{E} [(X_i - \nu_{X_i|Z})^2]}{\mathbb{E} [(X_i - \mu_{X_i|Z})^2]} \\
&= \frac{1}{2} \log \left(\frac{\mathbb{E} [(X_i - \nu_{X_i|Z} + \mu_{X_i|Z} - \mu_{X_i|Z})^2]}{\mathbb{E} [(X_i - \mu_{X_i|Z})^2]} \right) \\
&= \frac{1}{2} \log \left(\frac{\mathbb{E} [(X_i - \mu_{X_i|Z})^2] + \mathbb{E} [(\mu_{X_i|Z} - \nu_{X_i|Z})^2] + 2\mathbb{E} [(X_i - \mu_{X_i|Z})(\mu_{X_i|Z} - \nu_{X_i|Z})]}{\mathbb{E} [(X_i - \mu_{X_i|Z})^2]} \right) \\
&= \frac{1}{2} \log \left(1 + \frac{\mathbb{E} [(\mu_{X_i|Z} - \nu_{X_i|Z})^2] + 2\mathbb{E}_Z \mathbb{E}_{X_i|Z} [(X_i - \mu_{X_i|Z})(\mu_{X_i|Z} - \nu_{X_i|Z})]}{\mathbb{E} [(X_i - \mu_{X_i|Z})^2]} \right) \\
&= \frac{1}{2} \log \left(1 + \frac{\mathbb{E} [(\mu_{X_i|Z} - \nu_{X_i|Z})^2]}{\mathbb{E} [(X_i - \mu_{X_i|Z})^2]} \right) \geq 0.
\end{aligned}$$

We see that this regularizer is always non-negative and is zero exactly for modular latent factor models (when $\mu_{X_i|Z} = \nu_{X_i|Z}$). Summing up, the final objective simplifies to the following:

$$\underset{W}{\text{minimize}} \sum_{i=1}^p 1/2 \log \mathbb{E} [(X_i - \nu_{X_i|Z})^2] + \sum_{j=1}^m 1/2 \log \mathbb{E} [Z_j^2]. \quad (10)$$

This objective depends on pairwise statistics and requires no matrix inversion. The global minimum is achieved for modular latent factor models. The next step is to approximate the expectations in the objective (3) with empirical means and optimize it with respect to the parameters W .

After training the method we can interpret $\hat{\pi}_i \in \arg \max_j I(Z_j; X_i) = \arg \max_j -1/2 \log(1 - R_{j,i}^2) = \arg \max_j |R_{j,i}|$ as the parent of variable X_i . Additionally, we can estimate the covariance matrix of the observed variables. The method we use for estimating the covariance is as follows. First, we have assumed that the data is standardized, so we just need to calculate the off-diagonal terms. If $TC(X|Z) = 0$, this implies the conditional covariance of X given Z is diagonal. Additionally, using the law of total covariance we have:

$$\text{Cov}[X_i, X_{\ell \neq i}] = \mathbb{E} [\text{Cov}[X_i, X_{\ell}|Z]] + \text{Cov}[\mu_{X_i|Z}, \mu_{X_{\ell}|Z}].$$

By combining the last two statements we get:

$$\mathbb{E} [\text{Cov}[X_i, X_{\ell \neq i}|Z]] = \mathbb{E} [X_i X_{\ell}] - \mathbb{E} [\mu_{X_i|Z} \mu_{X_{\ell}|Z}] = 0.$$

If we assume the constraints $TC(Z) = 0$ & $\forall i, TC(Z|X_i) = 0$ are satisfied, we saw that this implies $\mu_{X_i|Z} = \nu_{X_i|Z}$. Also, as $TC(Z) = 0 \Rightarrow \mathbb{E} [Z_j Z_k] = \delta_{j,k} \mathbb{E} [Z_j^2]$, the off-diagonal elements of $\mathbb{E} [X_i X_{\ell}]$ satisfy:

$$\mathbb{E} [X_i X_{\ell \neq i}] = \mathbb{E} [\nu_{X_i|Z} \nu_{X_{\ell}|Z}] = \frac{(B^T B)_{i,\ell}}{(1+r_i)(1+r_{\ell})}.$$

In conclusion we get the following covariance matrix estimates:

$$\hat{\Sigma}_{i,\ell \neq i} = \frac{(B^T B)_{i,\ell}}{(1+r_i)(1+r_{\ell})}, \quad \hat{\Sigma}_{i,i} = 1. \quad (11)$$

Note that the covariance matrix estimate corresponds to the covariance matrix of the learned model if $TC(X|Z) = 0, TC(Z) = 0$, and $\forall i, TC(Z|X_i) = 0$, i.e. the learned model is modular. Otherwise it is an approximation of to the covariance matrix of the learned model. From Eq. 11 we see that the estimates are low-rank plus diagonal matrices. In case when the learned model is modular, it is also block-diagonal with each block being a diagonal plus rank-one matrix. Therefore, encouraging modular structures pushes the low-rank covariance estimate to be also block-diagonal with each block being a diagonal plus rank-one matrix.

B.1 Derivation of the conditional mean under modularity constraints

Under the conditions that X, Z are jointly Gaussian and $\forall i, TC(X|Z_i) = 0$, we would like to derive the mean of X_i conditioned on Z , $\nu_{X_i|Z}$. We have that $TC(Z|X_i) = 0 \Rightarrow p(x_i|z) = p(x_i)/p(z) \prod_j p(z_j|x_i)$. We will look at the distribution $q(x_i|z) = p(x_i)/p(z) \prod_j p(z_j|x_i)$ and calculate the conditional mean of this distribution.

Let $R_{j,i}$ be the Pearson correlation coefficient between Z_j and X_i whose means and standard deviations are respectively indicated with ν_j, ρ_j and μ_i, σ_i (all with respect to the distribution p). The marginal distribution for the Gaussian distribution relating Z_j and X_i is well known:

$$p(z_j|x_i) = \mathcal{N}(\nu_j + R_{j,i}\rho_j/\sigma_i(x_i - \mu_i), (1 - R_{j,i}^2)\rho_j^2).$$

Now we look only at the exponents of $q(x_i|z)$, ignoring the normalization, to get the following:

$$-\log q(x_i|z) \propto (x_i - \mu_i)^2/\sigma_i^2 + \sum_{j=1}^m (z_j - \nu_j - R_{j,i}\rho_j/\sigma_i(x_i - \mu_i))^2/((1 - R_{j,i}^2)\rho_j^2).$$

Collecting only the terms involving x_i we get the following:

$$\begin{aligned} -\log q(x_i|z) &\propto Ax_i^2 + Bx_i + C, \\ \text{with } A &= 1/\sigma_i^2 + \sum_{j=1}^m \frac{R_{j,i}^2\rho_j^2/\sigma_i^2}{(1 - R_{j,i}^2)\rho_j^2}, \quad B = -2\mu_i/\sigma_i^2 - \sum_{j=1}^m \frac{2(z_j - \nu_j + \mu_i R_{j,i}\rho_j/\sigma_i)R_{j,i}\rho_j/\sigma_i}{(1 - R_{j,i}^2)\rho_j^2}. \end{aligned}$$

From completing the square, we see that the conditional mean of $X_i|Z$ has the form $\nu_{X_i|Z} = -B/(2A)$.

Finally, we simplify the formulae because $\mu_i = \mathbb{E}[X_i] = \nu_j = \mathbb{E}[Z_j] = 0$ and $\sigma_i^2 = \mathbb{E}[X_i^2] = 1$.

This implies that $R_{j,i} = \mathbb{E}[X_i Z_j] / \sqrt{\mathbb{E}[X_i^2] \mathbb{E}[Z_j^2]}$, leaving us with the following form:

$$\nu_{X_i|Z} = \frac{1}{1 + r_i} \sum_{j=1}^m B_{j,i} \frac{Z_j}{\sqrt{\mathbb{E}[Z_j^2]}}, \quad \text{with } B_{j,i} = \frac{R_{j,i}}{1 - R_{j,i}^2}, r_i = \sum_{j=1}^m R_{j,i} B_{j,i}.$$

C Sample complexity lower bound

In this section we derive a lower bound on sample complexity for learning the structure of modular latent factor model. We follow the construction of information-theoretic sample complexity bounds in [7].

Theorem C.1. For a multivariate Gaussian modular latent factor model with p observed variables $X_{1:p}$, m latent variables $Z_{1:m}$ with p/m children each and additive white Gaussian noise channel from parent to child with signal-to-noise ratio s , the number of samples, n , required to recover the structure of the graphical model with error probability ϵ is lower bounded as

$$n \geq \frac{2 \left((1 - \epsilon) \log \left(\binom{p}{p/m, \dots, p/m} \frac{1}{m!} \right) - 1 \right)}{(p - 1) \log(1 + s \frac{1-1/m}{1-1/p}) - (m - 1) \log(1 + s \frac{p}{m})}. \quad (12)$$

Proof. Consider the class of modular latent factor models with p observed variables and m latent factors each having exactly p/m children. To distinguish the structure among this class of models corresponds to partitioning the observed variables into m equally sized groups. The number of such groupings is,

$$M = \binom{p}{p/m, \dots, p/m} \frac{1}{m!},$$

the multinomial coefficient for dividing p items into m equally sized boxes, divided by the number of indistinguishable permutations among boxes, $m!$. We take $\theta \in \{1, \dots, M\}$ to be an index specifying a model in this ensemble. Now learning the structure corresponds to finding θ from data.

W.l.o.g. assume $\forall j, \text{Var}[Z_j] = b$. Then $X_i = Z_{\pi_\theta(i)} + \eta_i$, where $\pi_\theta(i)$ is the index of the parent of X_i in model θ and η_i is independent Gaussian noise with variance a . Since we have fixed the signal-to-noise ratio, we have that $a = b/s$. W.l.o.g. we can assume that $\forall i, \mathbb{E}[X_i] = 0$. Then the covariance matrix of observed variables, $\Sigma_{\theta,i,j} = \mathbb{E}[X_i X_j] = b\delta_{\pi_\theta(i),\pi_\theta(j)} + a\delta_{i,j}$, where δ is the Kronecker delta.

Fano's inequality tells us that the probability of an error, ϵ , in picking the correct index θ given n samples of data, $X_{1:p}^{1:n}$, is bounded as follows:

$$\epsilon \geq 1 - \frac{I(\theta; X_{1:p}^{1:n}) + 1}{\log M}.$$

Following [7], we use an upper bound for the mutual information, $I(\theta; X_{1:p}^{1:n}) \leq nF/2$, where

$$F = \log \det \bar{\Sigma} - 1/M \sum_{\theta=1}^M \log \det \Sigma_\theta,$$

and $\bar{\Sigma} = 1/M \sum_{\theta=1}^M \Sigma_\theta$. Re-arranging Fano's inequality gives the following sample complexity bound:

$$n \geq 2 \frac{(1 - \epsilon) \log M - 1}{F}.$$

All that remains is to find an expression for F . To build intuition, we explicitly write out the case for $p = 4, m = 2$, and for some θ .

$$\Sigma_\theta = \begin{bmatrix} b+a & b & 0 & 0 \\ b & b+a & 0 & 0 \\ 0 & 0 & b+a & b \\ 0 & 0 & b & b+a \end{bmatrix}$$

Clearly this is a block diagonal matrix where each block is a diagonal plus rank-one (DPR1) matrix. After we average over all θ to get $\bar{\Sigma}$, every off-diagonal entry will be the same, equal to the probability of $j \neq i$ being in the same group as i , or $(p/m - 1)/(p - 1)$. Therefore $\bar{\Sigma}$ is also a DPR1 matrix. Using standard identities for block diagonal and DPR1 matrices, we calculate the determinants:

$$\det \Sigma_\theta = a^p \left(1 + \frac{b}{a} \frac{p}{m}\right)^m,$$

$$\det \bar{\Sigma} = a^p \left(1 + \frac{b}{a} \frac{p}{m}\right) \left(1 + \frac{b}{a} \frac{p}{m} \left(\frac{m-1}{p-1}\right)\right)^{p-1}.$$

Finally, we can combine all of these expressions to get a lower bound for sample complexity that depends only on p, m , and the signal-to-noise ratio, $s = b/a$. \square

The bound of Thm. [C.1] is not very intuitive because it involves logarithm of a multinomial coefficient. We provide a simpler asymptotic expression for the bound. Using Stirling's approximation we have that $\log \binom{p}{p/m, \dots, p/m} \frac{1}{m!} \approx p \log m + 1/2 \log(p/m) - m/2 \log(m p 2\pi/e^2)$ for large values of p . In the limit of large p , this approximation gives us the following lower bound:

$$n \geq \frac{2(1 - \epsilon) \log m}{\log(1 + s(1 - 1/m))}.$$

We see that in the limit of large p the bound becomes constant rather than becoming infinite. Moreover, when we plot the lower bound of Eq. [12] in Fig. [7] we see that for fixed number of latent factors the bound goes down as we increase p . These two facts together hint (but do not prove) that modular latent factor models may allow blessing of dimensionality. An evidence of blessing of dimensionality is demonstrated in Sec. [4]. Intuitively, recovery gets easier because more variables provide more signal to reconstruct the fixed number of latent factors. While it is tempting to retrospectively see this as obvious, the same argument could be (mistakenly) applied to other families of latent factor models, such as the unconstrained latent factor models shown in Fig. [1a] for which the sample complexity grows as we increase p [2, 15].

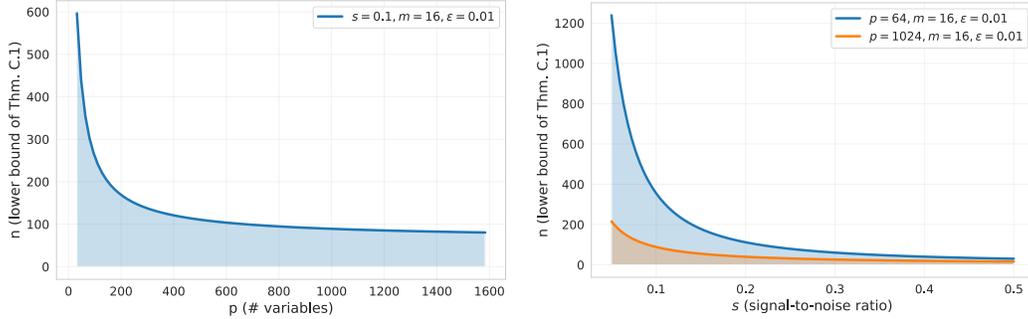


Figure 7: Theorem [C.1](#) prevents perfect structure recovery in the shaded region. On the left: for fixed signal-to-noise ratio and number of latent factors, the lower bound of Thm. [C.1](#) decreases as the number of observed variables increases. On the right: the same effect is visible for other values of signal-to-noise ratio.

D Implementation details

In this section we present details on baselines, experiments, and hyperparameters.

Baselines For factor analysis, PCA, sparse PCA, independent component analysis, k-means clustering, spectral clustering, negative matrix factorization, hierarchical agglomerative clustering using Euclidean distance, hierarchical agglomerative clustering the Ward linkage rule, Ledoit-Wolf and graphical LASSO we used the scikit-learn implementations [\[42\]](#). We implemented latent tree modeling with the “Relaxed RG” method. We slightly modified latent tree modeling to use the same prior information as other methods in the comparison, namely, that there are exactly m groups and observed nodes can be siblings, but not parent and child. For latent variable graphical LASSO, we used the implementation available in the REGAIN repository [\[3\]](#)

Experimental setups In the blessing of dimensionality experiments all methods were given the correct number of clusters. The scores were computed using 10000 test examples. When possible we reported the means and standard deviations over 20 runs. In the covariance estimation experiments with synthetic data, models requiring a number of latent factors or a number of components were given the correct number. The scores were computed using 1000 test examples. We reported the means and standard deviations over 5 runs. In the stock market experiments models were trained on n weeks and their estimates were evaluated using the negative log-likelihood on the subsequent 26 weeks. We presented the average score from rolling the training and testing sets over the entire time period. Standard deviations are not presented because scores corresponding to different time periods are very different, resulting in large standard deviations. This is due to the stock market exhibiting different behaviour in different time periods. In experiments with OpenML datasets we used a random 80-20 train-test split. We reported the negative log-likelihood on test sets. As large amount of computation is needed to generate results on OpenML datasets, we did only a single run for each dataset.

Hyperparameters In all cases the proposed method was trained using Adam optimizer with 0.01 learning rate, $\beta_1 = 0.9$, and $\beta_2 = 0.999$. In all covariance estimation problems the hyperparameters were selected from a grid of values using a 3-fold cross-validation procedure. The sparsity parameter of sparse PCA was selected from $[0.1, 0.3, 1.0, 3.0, 10.0]$. The sparsity parameters of GLASSO and latent variable GLASSO were selected from $[0.01, 0.1, 0.3, 1.0, 3.0, 10.0]$. For latent variable GLASSO, the additional regularization parameter (“tau”, controlling the nuclear norm of the low-rank part of the inverse covariance matrix) was selected from $[0.01, 0.1, 1.0, 10.0, 100.0]$. In the experiments with OpenML datasets the sparsity hyperparameter of BigQUIC was selected from $[2^0, 2^1, 2^2, 2^3]$. In the timing experiments the sparsity parameters of sparse PCA and GLASSO were set to 1.0. LVGLASSO was trained with the sparsity parameter set to 0.1 and with “tau” set to 30.0.

<https://github.com/fdtomasi/regain>

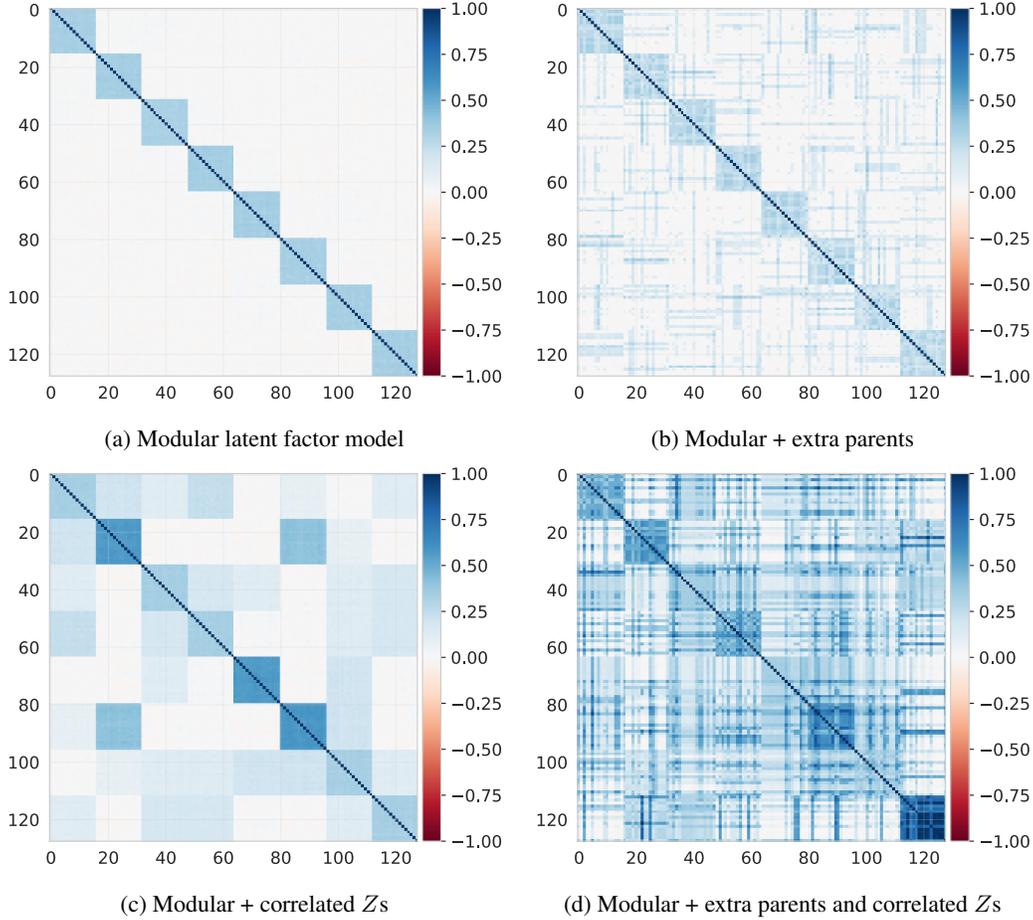


Figure 8: Empirical covariance matrices (estimated using $n = 10^4$ samples) corresponding to modular (a) and approximately modular (b, c, d) latent factor models. In all examples $m = 8$, $p = 128$, $s = 0.5$.

E Details on generating synthetic data

In all experiments involving a synthetic modular latent factor model we generate the data the following way. We first take m independent standard Gaussian random variables, $Z_1, Z_2, \dots, Z_m \stackrel{iid}{\sim} \mathcal{N}(0, 1)$. For simplicity we assume that m divides p and each latent factor has exactly p/m children. W.l.o.g. we connect the first p/m observed variables with Z_1 , then next p/m variables with Z_2 and so on. We assume additive white Gaussian noise channel with signal-to-noise ratio s from each parent to its children. In this setup, we set $X_i = \sqrt{\frac{s}{s+1}} Z_{\pi_i} + \sqrt{\frac{1}{s+1}} \eta_i$, where π_i is the index of the parent of X_i , and η_i is independent standard Gaussian noise. Fig. 8a shows a covariance matrix corresponding to a modular latent factor models created using the described procedure.

To create approximately modular latent factor models we do two modifications on a modular latent factor model: correlating the latent variables and adding extra parents for observed variables. For correlating the latent factors we take m independent standard normal random variables $\xi_j, j = 1..m$ and compute $z_j = (\sqrt{2}\xi_j + \xi_u + \xi_v)/2$, where $u, v \sim \text{Uniform}\{1, 2, \dots, m\}$. For adding extra parents, we randomly sample p extra edges from a latent factor to a non-child observed variable. By this we create on average one extra edge per each observed variable. To keep the notion of clusters well-defined, we make sure that each observed variable has higher mutual information with its main parent compared to that with added extra parents. Suppose some X_i has k extra parents, $Z_{\tau_1}, \dots, Z_{\tau_k}$. Then we splits $\frac{s}{s+1}$ – the variance of the signal in a pure modular case – into $k + 2$ equal parts,

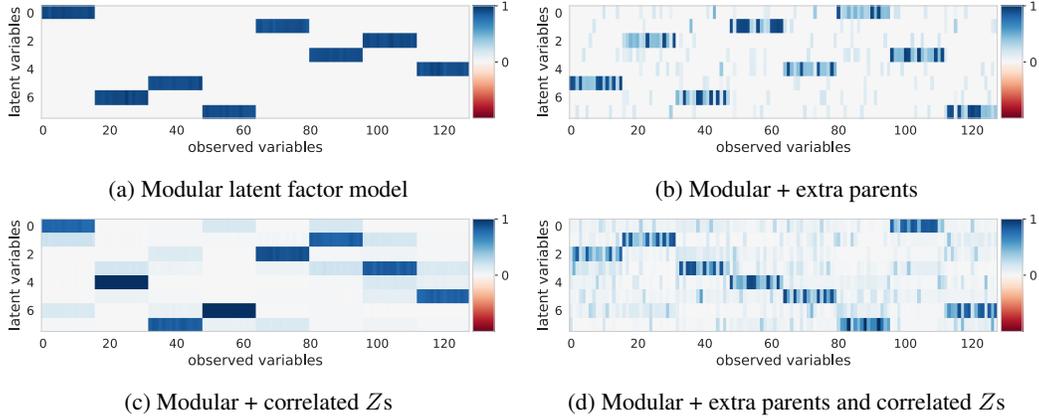


Figure 9: Mutual information matrices between observed variables and latent factors linear CorEx produces when it is trained on a modular (a) and approximately modular (b, c, d) latent factor models. In all examples $m = 8$, $p = 128$, $s = 5.0$.

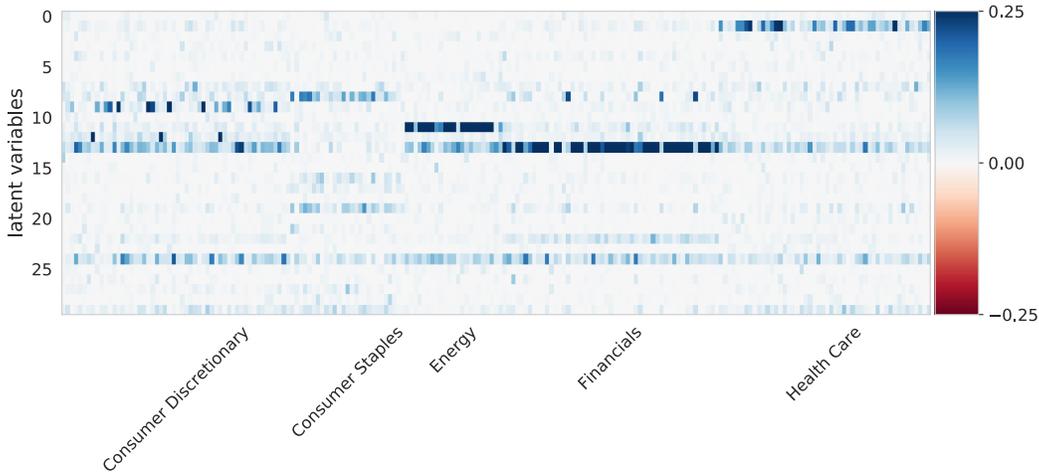


Figure 10: Mutual information matrix between observed variables (stocks) and latent factors linear CorEx produces when trained on the stock marked data (January 2014-January 2017).

$\delta = \frac{s}{(s+1)(k+2)}$. We then set $X_i = \sqrt{2\delta}Z_{\pi_i} + \sqrt{\delta}Z_{\tau_1} + \dots + \sqrt{\delta}Z_{\tau_k} + \sqrt{\frac{1}{s+1}}\eta_i$, where again η_i is independent standard Gaussian noise. Figs. 8b, 8c, 8d show covariance matrices corresponding to approximately modular latent factor models created using the described procedures.

F Additional results

In this section we provide additional results that were not presented in the main text due to the space constraints.

F.1 Examining the modularity of learned models

We do visualizations to see whether the regularization term of linear CorEx actually leads to learning modular (or approximately modular) latent factor models. We examine the mutual information matrices between observed and latent variables that linear CorEx produces when it is trained on different types of synthetic data (see Fig 9). We see that the regularization term we add for encouraging modular structures is indeed effective.

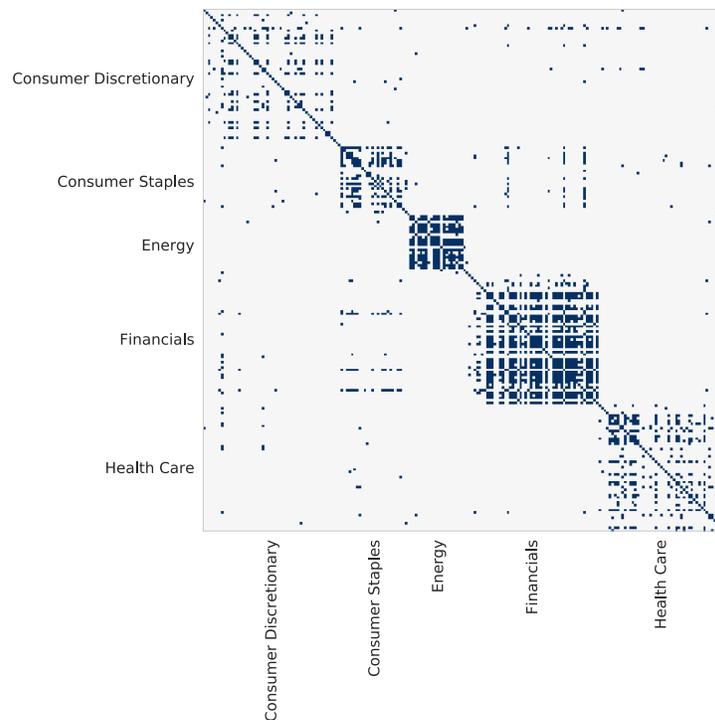


Figure 11: Inverse covariance matrix of some of the S&P 500 stocks. Plotted are the cells that have absolute value greater than 0.015.

Next, we look at the same mutual information matrix for stock market data. Fig. 10 shows the mutual information matrix for S&P 500 stocks belonging to “consumer discretionary”, “consumer staples”, “energy”, “financials”, and “health care” sectors. We see that most of the stocks have significant mutual information only with a few latent factors. Moreover, stocks belonging to the same sector are likely to share a parent. Additionally, we visualize the inverse covariance matrix of these stocks (see Fig. 11). For Gaussian random variables the thresholded inverse covariance matrix can be interpreted as a random Markov field. We see that it is almost block-diagonal, but has some off-diagonal connections, confirming that the learned model is close to being a modular latent factor model.

Summing up, all these visualizations assert that the linear CorEx succeeds in biasing the model selection procedure towards modular structures. More importantly, we see that when the pure modular structure is inappropriate, it picks solutions that are close to being modular.

F.2 Results on OpenML datasets

Table 2 presents a comparison of various covariance estimation baselines on 51 OpenML datasets.

Table 2: This table compares covariance estimates on OpenML data. Scores reported are negative log-likelihood (lower better) and the best entry is bolded. Scores orders of magnitude larger than the best score or evaluating to NaN are shortened with “*”. Methods compared, in order, are PCA, Sparse PCA, factor analysis, Ledoit-Wolf, GLASSO (using the BigQUIC algorithm), and the method proposed in this paper.

ID:Dataset	p	n	Methods					
			PCA	SPCA	FA	LW	BigQUIC	Proposed
5:arrhythmia	206	54	178	-33	*	164	*	-74
407:krystek	1143	24	2122	-1748	*	707	-1428	-2816
408:depreux	1143	20	1454	*	*	852	*	-482
409:pdgfr	321	63	112	40	*	83	364	-6
410:carbolenes	1143	29	1900	*	*	923	*	-8
419:PHENETYLI	629	17	876	560	5584	281	1041	286
420:cristalli	1143	25	1846	*	*	1366	*	780
424:pah	113	64	-129	47	-188	25	134	58
439:chang	1143	27	2331	*	*	1058	*	-6
1017:arrhythmia	206	54	178	-33	*	164	*	-60
1104:leukemia	7129	57	17028	13164	396636	7019	11530	7336
1107:tumorsC	7129	48	16990	8499	9642	8070	9398	8399
1122:APBreastProstat	10935	330	18427	17219	17741	13431	17002	10639
1123:APEndometriumBr	10935	324	18960	12616	12720	11356	18330	10452
1124:APOmentumUterus	10935	160	84928	82496	82656	66784	76832	66176
1125:APOmentumProsta	10935	116	90024	100392	100032	69168	81264	67560
1126:APColonLung	10935	329	84612	76362	76626	66198	76098	67188
1127:APBreastOmentum	10935	336	86020	88196	88604	66824	79968	68408
1128:OVABreast	10935	1236	83626	79434	79087	64951	76483	70308
1129:APUterusKidney	10935	307	85498	74276	74214	68882	75764	68882
1130:OVALung	10935	1236	81989	*	73904	81518	76409	69291
1131:APP prostateUteru	10935	154	85653	73687	73625	67208	74617	67363
1132:APOmentumLung	10935	162	88803	83820	83424	70917	79002	68805
1133:APEndometriumCo	10935	277	84840	75376	75152	65576	76776	66920
1134:OVAKidney	10935	1236	81964	75144	73507	81592	76210	69242
1135:APColonProstate	10935	284	84702	82365	82194	65550	78318	67260
1136:APLungUterus	10935	200	87200	77880	77360	69200	76480	68440
1137:APColonKidney	10935	436	83776	73515	73084	66616	75882	68094
1138:OVAUterus	10935	1236	81964	74772	73358	81493	76136	69242
1139:OVAOmentum	10935	1236	81964	75442	74152	81567	76458	69266
1140:APOvaryLung	10935	259	87464	*	85280	86320	79560	68380
1141:APEndometriumPr	10935	104	90006	85449	84063	69447	77994	67263
1142:OVAEndometrium	10935	1236	81989	74574	73086	81493	76062	69242
1143:APColonOmentum	10935	290	84332	77198	77140	65250	76908	66816
1144:APP prostateKidne	10935	263	85277	79553	79553	69589	77857	68052
1145:APBreastColon	10935	504	84416	77346	77305	65165	78144	68448
1146:OVAProstate	10935	1236	81989	75318	73160	81542	76434	69266
1147:APOmentumKidney	10935	269	84780	75114	74952	68796	77112	67770
1148:APBreastUterus	10935	374	85500	73275	73155	66892	79575	68475
1149:APOvaryKidney	10935	366	86062	*	76590	85470	77626	68835
1150:APBreastLung	10935	376	86868	86032	86260	68195	80332	69441
1151:APEndometriumOm	10935	110	89474	98868	100386	68178	80388	66550
1152:APP prostateOvary	10935	213	87118	*	75594	86473	78045	68026
1153:APColonOvary	10935	387	85488	*	91572	85254	81198	68047
1154:APEndometriumLu	10935	149	89490	74940	75180	70350	76710	67260
1233:eating	6373	756	7843	6703	5381	-1110	7980	5457
1457:amazon-commerce	10000	1200	15576	11376	11256	10680	12216	10920
1458:arcene	10000	160	19181	9152	8746	-1267	*	8179
1484:lsvt	310	100	200	152	872	212	464	180
1514:micro-mass	1300	288	1166	50547	50912	1056	*	-708
1515:micro-mass	1300	456	1260	8041	71493	1224	*	589
Total # wins			0	1	1	18	0	32