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On the Use of Minimum Penalties in Statistical Learning

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Abstract

Modern multivariate machine learning and statistical methodologies estimate parameters of interest while leveraging prior knowledge of the association between outcome variables. The methods that do allow for estimation of relationships do so typically through an error covariance matrix in multivariate regression which does not scale to other types of models. In this article we proposed the MinPEN framework to simultaneously estimate regression coefficients associated with the multivariate regression model and the relationships between outcome variables using mild assumptions. The MinPen framework utilizes a novel penalty based on the minimum function to exploit detected relationships between responses. An iterative algorithm that generalizes current state of the art methods is proposed as a solution to the nonconvex optimization that is required to obtain estimates. Theoretical results such as high dimensional convergence rates, model selection consistency, and a framework for post selection inference are provided. We extend the proposed MinPen framework

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to other exponential family loss functions, with a specific focus on multiple binomial responses. Tuning parameter selection is also addressed. Finally, simulations and two data examples are presented to show the finite sample properties of this framework.

Keywords: Non-Convex Optimization, Post-Selection Inference, High Dimensional Convergence, Graph Constrained Models, Selection Consistency

1 Introduction

Multivariate, also known as multiple response or multi-task, regression jointly models multiple responses (outcomes) given a common set of predictors (inputs). Joint modeling of responses, as opposed to modeling each response separately, is only interesting if the responses are related. Leveraging these relationships should improve estimation of the model, but in practice these relationships may not always be known. In this work we propose a framework to simultaneously estimate the regression coefficients and detect relationships between the response variables. The framework also allows for simultaneous variable selection.

Consider the sample of independent data, $\{\mathbf{x}_i, \mathbf{y}_i\}_{i=1}^n$, where $\mathbf{x}_i \in \mathbb{R}^p$ and $\mathbf{y}_i \in \mathbb{R}^r$ with a model of

$$\mathbf{y}_i = \mathbf{B}^{*T} \mathbf{x}_i + \boldsymbol{\epsilon}_i, \tag{1}$$

where $\mathbf{B}^* = (\boldsymbol{\beta}_1^*, \dots, \boldsymbol{\beta}_r^*) \in \mathbb{R}^{p \times r}$ and $\boldsymbol{\epsilon}_i \in \mathbb{R}^r$ are i.i.d. random vectors with mean zero and covariance matrix $\boldsymbol{\Sigma}$. A popular estimator for \mathbf{B}^* is

$$\widehat{\mathbf{B}} = \underset{\boldsymbol{\beta}_{1},\dots,\boldsymbol{\beta}_{r}\in\mathbb{R}^{p}}{\arg\min} \frac{1}{2n} \sum_{k=1}^{r} \sum_{i=1}^{n} (y_{ik} - \mathbf{x}_{i}^{T} \boldsymbol{\beta}_{k})^{2},$$
(2)

which has desirable qualities such as consistency and asymptotic normality under mild conditions, and is the maximum likelihood estimator if the errors are from a multivariate normal distribution. However, there is something unsatisfying about not using correlation of the responses in the estimation. To improve efficiency in estimation of coefficients, while accounting for correlation across the errors Rothman et al. (2010) (MRCE), Witten & Tibshirani (2009) (SCOUT) and Lee & Liu (2012) have all proposed methods which simultaneously estimates the regression coefficients and the inverse covariance matrix of the errors. These methods all aim to improve estimation of regression coefficients by exploiting correlation across the responses that is not explained by the predictors.

We propose an alternative approach that focuses on relationships across the responses that can be explained, due to similarity in the regression coefficients. Specifically, we consider the kth and lth response related if β_k^* is similar to β_l^* or $-\beta_l^*$. If relationships were known *a priori*, then for each response *k* one could define three disjoint sets: (1) P_k , a set of responses positively related to response *k*; (2) N_k , a set of responses negatively related to response *k*; and (3) Z_k a set of responses that are not related to *k*. These relationships could be used to improve estimation using a penalized likelihood such as

$$\underset{\mathbf{B}\in\mathbb{R}^{p\times r}}{\operatorname{arg\,min}} \frac{1}{2n} \sum_{k=1}^{r} (\mathbf{y}_{k} - \mathbf{x}_{i}^{T}\boldsymbol{\beta}_{k})^{T} (\mathbf{y}_{k} - \mathbf{x}_{i}^{T}\boldsymbol{\beta}_{k}) + \frac{\gamma}{2} \sum_{k=1}^{r} \left(\sum_{l\in P_{k}} \|\boldsymbol{\beta}_{l} - \boldsymbol{\beta}_{k}\|_{2}^{2} + \sum_{m\in N_{k}} \|\boldsymbol{\beta}_{m} + \boldsymbol{\beta}_{k}\|_{2}^{2} + \sum_{\substack{s\in Z_{k} \\ (3)}} \|\boldsymbol{\beta}_{k}\|_{2}^{2} \right) + \frac{\gamma}{2} \sum_{k=1}^{r} \left(\sum_{l\in P_{k}} \|\boldsymbol{\beta}_{l} - \boldsymbol{\beta}_{k}\|_{2}^{2} + \sum_{m\in N_{k}} \|\boldsymbol{\beta}_{m} + \boldsymbol{\beta}_{k}\|_{2}^{2} + \sum_{\substack{s\in Z_{k} \\ (3)}} \|\boldsymbol{\beta}_{m} - \boldsymbol{\beta}_{k}\|_{2}^{2} \right) + \frac{\gamma}{2} \sum_{k=1}^{r} \left(\sum_{l\in P_{k}} \|\boldsymbol{\beta}_{l} - \boldsymbol{\beta}_{k}\|_{2}^{2} + \sum_{m\in N_{k}} \|\boldsymbol{\beta}_{m} - \boldsymbol{\beta}_{k}\|_{2}^{2} + \sum_{\substack{s\in Z_{k} \\ (3)}} \|\boldsymbol{\beta}_{m} - \boldsymbol{\beta}_{k}\|_{2}^{2} \right) + \frac{\gamma}{2} \sum_{k=1}^{r} \left(\sum_{l\in P_{k}} \|\boldsymbol{\beta}_{l} - \boldsymbol{\beta}_{k}\|_{2}^{2} + \sum_{m\in N_{k}} \|\boldsymbol{\beta}_{m} - \boldsymbol{\beta}_{m}\|_{2}^{2} + \sum_{\substack{s\in Z_{k} \\ (3)}} \|\boldsymbol{\beta}_{m} - \boldsymbol{\beta}_{k}\|_{2}^{2} \right) + \frac{\gamma}{2} \sum_{k=1}^{r} \left(\sum_{l\in P_{k}} \|\boldsymbol{\beta}_{l} - \boldsymbol{\beta}_{k}\|_{2}^{2} + \sum_{m\in N_{k}} \|\boldsymbol{\beta}_{m} - \boldsymbol{\beta}_{m}\|_{2}^{2} \right) + \frac{\gamma}{2} \sum_{k=1}^{r} \left(\sum_{l\in P_{k}} \|\boldsymbol{\beta}_{l} - \boldsymbol{\beta}_{k}\|_{2}^{2} + \sum_{l\in P_{k}} \|\boldsymbol{\beta}_{l} - \boldsymbol{\beta}_{l}\|_{2}^{2} \right) + \frac{\gamma}{2} \sum_{k=1}^{r} \left(\sum_{l\in P_{k}} \|\boldsymbol{\beta}_{l} - \boldsymbol{\beta}_{k}\|_{2}^{2} + \sum_{l\in P_{k}} \|\boldsymbol{\beta}_{l} - \boldsymbol{\beta}_{l}\|_{2}^{2} \right) + \frac{\gamma}{2} \sum_{k=1}^{r} \left(\sum_{l\in P_{k}} \|\boldsymbol{\beta}_{l} - \boldsymbol{\beta}_{k}\|_{2}^{2} \right) + \frac{\gamma}{2} \sum_{k=1}^{r} \left(\sum_{l\in P_{k}} \|\boldsymbol{\beta}_{l} - \boldsymbol{\beta}_{l}\|_{2}^{2} \right) + \frac{\gamma}{2} \sum_{k=1}^{r} \left(\sum_{l\in P_{k}} \|\boldsymbol{\beta}_{l} - \boldsymbol{\beta}_{l}\|_{2}^{2} \right) + \frac{\gamma}{2} \sum_{k=1}^{r} \left(\sum_{l\in P_{k}} \|\boldsymbol{\beta}_{l} - \boldsymbol{\beta}_{l}\|_{2}^{2} \right) + \frac{\gamma}{2} \sum_{k=1}^{r} \left(\sum_{l\in P_{k}} \|\boldsymbol{\beta}_{l} - \boldsymbol{\beta}_{l}\|_{2}^{2} \right) + \frac{\gamma}{2} \sum_{k=1}^{r} \left(\sum_{l\in P_{k}} \|\boldsymbol{\beta}_{l}\|_{2}^{2} \right) + \frac{\gamma}{$$

where, for a vector $a \in \mathbb{R}^d$ define $||a||_q = (\sum_{j=1}^d a_j^q)^{\frac{1}{q}}$ as the L_q norm and $\gamma > 0$ is a tuning parameter to promote similarities between groups of related coefficients.

The method we propose is a generalization of (3) when the sets P_k , N_k and Z_k are not known *a priori* and uses an L_1 penalty for feature selection. The proposed method simultaneously estimates these unknown sets, while using the framework of (3) to include these detected relationships in the estimation. To simultaneously estimate the group structure we propose a novel penalty based on the minimum function. The penalized objective function is non-convex but can be re-stated as solving a fixed number of convex problems. However, this quickly becomes computationally intractable for even small values of r. Unlike the earlier cited work which focuses on structure due to relationships in the errors, this approach can be easily generalizable to simultaneous modeling of binary response variables or other non-continuous values by replacing the least squares loss function with the appropriate generalized linear model loss function.

In the least squares case we provide rates of convergence, model selection consistency and post selection inference results which hold for high-dimensional predictors, p >> n. The results hold for any grouping of the response variables and thus do not depend on the relationships between responses to be estimated correctly. To derive rates of convergence under standard conditions we generalize the results of Negahban et al. (2012), which do not directly apply because of the ridge fusion penalty used in this framework.

Our proposal builds on methods which simultaneously estimate clusters and univariate regression models (Witten et al. 2014), multivariate regression models (Price & Sherwood 2018), and precision matrices (Price, Molstad & Sherwood 2021). All of which propose iterative algorithms that alternate between estimating the clustering structure, using k-means, and the model of interest. These approaches only accommodate positive relationships and rely on k-means, which provides distinct sets of clusters and may be unreliable for high-dimensional problems. The proposed method allows for more complex structures in the response variables and provides a global minimum at each iteration.

Extensive work has been done on exploiting relationships in multivariate regression to improve efficiency. For a recent survey of the field see Price, Allenbrand & Sherwood (2021) and citations within. Reduced rank regression reduces the dimension of the problem by constraining the rank of the coefficient matrix **B** while minimizing the multivariate least squares objective function to find a set of latent variables that increase prediction accuracy (Anderson et al. 1951, Velu & Reinsel 2013, Chen & Huang 2016). Various group lasso penalties have been used to promote structure across response or a combination of response and predictors. (Kim et al. 2012, Li et al. 2015) Instead of the proposed approach of simultaneously estimating relationships and coefficients, many have used a two-stage clustering approach. In the first stage the clusters are estimated and in the second stage the structure is incorporated into the estimation of the coefficients. Examples include hierarchical clustering for tree guided lasso (Kim et al. 2012), convex clustering in conjoint analysis (Chen et al. 2016), separating global and response specific features (Xu et al. 2015) and replacing high dimensional responses or predictors with clusters (Bühlmann et al. 2013, Zhou et al. 2017).

Other methods in univariate regression, such as the GRACE method, utilize known relationships, or relationships that are estimated *a priori*, to assist in increasing accuracy and select relevant variables (Li & Li 2008, 2010*b*). Others such as Zhao & Shojaie (2016) have investigated statistical inference frameworks for univariate graph constrained models, that also accommodate error in the estimation of the graph Laplacian.

The key difference between the proposed method and state of the art methods in the literature, is that our method simultaneously detects relationships between the responses, regardless of sign, and estimates regression coefficients without needing to estimate a co-variance matrix of the errors. In Section 2 we introduce the MinPen framework with the least squares loss function along with theoretical results and the proposed algorithm. In Section 3 we extend the framework to multiple binomial responses with discussion of how it may generalized to other exponential family based loss functions. Sections 4 and 5 present simulations to investigate finite sample properties of the proposed methods. Finally we present examples of our methodology in applications in genomics and substance abuse overdoses in Section 6.

2 Least Squares Model

2.1 Method

We consider estimating \mathbf{B}^* in (1) when there may be similarities in the mean functions of the different responses. Define,

$$P(\boldsymbol{\beta}_l, \boldsymbol{\beta}_k) = \min(\|\boldsymbol{\beta}_l - \boldsymbol{\beta}_k\|_2^2, \|\boldsymbol{\beta}_l + \boldsymbol{\beta}_k\|_2^2, \|\boldsymbol{\beta}_l\|_2^2).$$
(4)

If $P(\boldsymbol{\beta}_{l}^{*}, \boldsymbol{\beta}_{k}^{*}) = \|\boldsymbol{\beta}_{l}^{*} - \boldsymbol{\beta}_{k}^{*}\|_{2}^{2}$ this implies that the mean functions for response k and l are positively related. If $P(\boldsymbol{\beta}_{l}^{*}, \boldsymbol{\beta}_{k}^{*}) = \|\boldsymbol{\beta}_{l}^{*} + \boldsymbol{\beta}_{k}^{*}\|_{2}^{2}$ this implies negative relationship between the two mean functions, while $P(\boldsymbol{\beta}_{l}^{*}, \boldsymbol{\beta}_{k}^{*}) = \|\boldsymbol{\beta}_{l}^{*}\|_{2}^{2}$ implies little or no relationship between the mean functions. Therefore a penalty based on the minimum function can be used to simultaneously identify and leverage relationships between mean functions. Motivated by this, we propose the minimum penalty elastic net multivariate regression (MinPen) estimator as the solution to

$$\underset{\mathbf{B}\in\mathbb{R}^{p\times r}}{\arg\min}\frac{1}{2n}\sum_{i=1}^{n}\sum_{k=1}^{r}(y_{ik}-\mathbf{x}_{i}^{T}\boldsymbol{\beta}_{k})^{2}+\delta\sum_{k=1}^{r}\|\boldsymbol{\beta}_{j}\|_{1}+\frac{\gamma}{2}\sum_{l=1}^{r}\sum_{k=1,k\neq l}^{r}P(\boldsymbol{\beta}_{l},\boldsymbol{\beta}_{k}),$$
(5)

where δ and γ are non-negative tuning parameters specified by the user. We refer to this as an elastic net estimator as it uses the combination of the lasso and ridge penalties as first introduced by Zou & Hastie (2005). The lasso penalty, associated with tuning parameter δ , is used to simultaneously perform variable selection and estimate regression coefficients. While, the proposed ridge type penalty, associated with γ , is a non-convex penalty that simultaneously identifies and exploits relationships between coefficient vectors of different responses.

Both our theory and algorithms will deal with a vectorized version of the solution to (5). For a matrix \mathbf{A} , define vec(\mathbf{A}) as the vector formed by stacking the columns of \mathbf{A}

on top of one another. Define $\mathbf{Y} = (\mathbf{y}_1, \dots, \mathbf{y}_n)^T \in \mathbb{R}^{n \times r}, \ \mathbf{X} = (\mathbf{x}_1, \dots, \mathbf{x}_n)^T \in \mathbb{R}^{n \times r},$ $\mathbf{E} = (\epsilon_1, \dots, \epsilon_n)^T \in \mathbb{R}^{n \times r}, \ \tilde{\mathbf{y}} = \operatorname{vec}(\mathbf{Y}) \in \mathbb{R}^{nr}, \ \tilde{\mathbf{X}} = \mathbf{I}_r \otimes \mathbf{X} \in \mathbb{R}^{nr \times pr}, \ \boldsymbol{\beta}^* = \operatorname{vec}(\mathbf{B}^*) \in \mathbb{R}^{pr},$ $\tilde{\boldsymbol{\epsilon}} = \operatorname{vec}(\mathbf{E}) \in \mathbb{R}^{nr} \text{ and } \ \tilde{\boldsymbol{\Sigma}} = \boldsymbol{\Sigma} \otimes \mathbf{I}_n \in \mathbb{R}^{nr \times nr}.$ Then the vectorized version of (1) is

$$\tilde{\mathbf{y}} = \tilde{\mathbf{X}}\boldsymbol{\beta}^* + \tilde{\boldsymbol{\epsilon}}.$$
(6)

Define the set

$$\mathcal{A} = \left\{ \mathbf{A} \in \{-1, 0, 1\}^{r(r-1)p \times rp} \middle| ||\mathbf{A}\boldsymbol{\beta}||_2^2 = \sum_{l=1}^r \sum_{m=1, l \neq m}^r ||\boldsymbol{\beta}_l - d_{lm}\boldsymbol{\beta}_m||_2^2 \text{ where } d_{lm} \in \{-1, 0, 1\} \right\}.$$

The estimator in (5) is equivalent to

$$(\hat{\boldsymbol{\beta}}, \hat{\mathbf{A}}) = \operatorname*{argmin}_{\boldsymbol{\beta} \in \mathbb{R}^{pr}} \min_{\mathbf{A} \in \mathcal{A}} \left(\frac{1}{2n} || \tilde{\mathbf{Y}} - \tilde{\mathbf{X}} \boldsymbol{\beta} ||_{2}^{2} + \delta || \boldsymbol{\beta} ||_{1} + \frac{\gamma}{2} || \mathbf{A} \boldsymbol{\beta} ||_{2}^{2} \right).$$
(7)

Due to the minimum function the objective function is non-convex, which typically makes theoretical study of the global minimizer challenging. However, the global solution to (7) is the pair $(\hat{\boldsymbol{\beta}}, \hat{\mathbf{A}})$ that minimizes across the $|\mathcal{A}|$ potential convex objective functions. So while the objective function is nonconvex, the solution $\hat{\boldsymbol{\beta}}$ minimizes a convex function. If the matrix \mathbf{A} is known *a priori* the estimator of $\boldsymbol{\beta}^*$ is

$$\hat{\boldsymbol{\beta}}(\mathbf{A}) = \operatorname*{arg\,min}_{\boldsymbol{\beta} \in \mathbb{R}^{pr}} \frac{1}{2n} \sum_{i=1}^{n} \sum_{k=1}^{r} (y_i - \mathbf{x}_i^T \boldsymbol{\beta}_k)^2 + \delta \sum_{k=1}^{r} \|\boldsymbol{\beta}_k\|_1 + \frac{\gamma}{2n} \boldsymbol{\beta}^T \mathbf{A}^T \mathbf{A} \boldsymbol{\beta}.$$
(8)

For our theoretical results we first analyze the estimator $\hat{\boldsymbol{\beta}}(\mathbf{A})$. Then, using that the solution is one of the potential $|\mathcal{A}|$ solutions, we provide rates of convergence for the global solution, $\hat{\boldsymbol{\beta}}$. It may be desirable to replace \mathcal{A} with some subset, say $\mathcal{A}_1 \subset \mathcal{A}$. If it is known that two responses are positively related then it does not makes sense to consider the negative or no relationship for those two responses and thus a suitable \mathcal{A}_1 could be used instead of \mathcal{A} . For this paper we consider the general case, but the results and algorithm presented here will hold if a smaller set is considered.

2.2 Theoretical Results

Mild or standard conditions were used to derive the rates of convergence. Before presenting the conditions, we define some notation that is used in our theorems and conditions. The subspace for the active predictors is defined as $\mathcal{M} \equiv \{\mathbf{a} \in \mathbb{R}^{pr} | a_j = 0 \text{ if } \boldsymbol{\beta}_j^* = 0\}$, with cardinality $s = |\mathcal{M}|$. The parameter space is separated using projections of vectors into orthogonal complements. Define $\mathbf{u}_{\mathcal{M}}$ as a projection of a vector \mathbf{u} into space $\mathcal{M}, \mathcal{M}^{\perp}$ as the orthogonal complement of \mathcal{M} and $\mathcal{C} \equiv \{\mathbf{a} \in \mathbb{R}^{pr} | || \mathbf{a}_{\mathcal{M}^{\perp}} ||_1 \leq 3 || \mathbf{a}_{\mathcal{M}} ||_1 \}$.

Condition 1. Define $\mathbf{x}_j \in \mathbb{R}^p$ to be the *j*th column vector of \mathbf{X} . For all $j \in \{1, \ldots, p\}$ $\frac{||\mathbf{x}_j||_2^2}{n} \leq 1.$

Condition 2. There exists positive constants κ_l and κ_u such that for all $\boldsymbol{\theta} \in C$ that $\kappa_l \leq \boldsymbol{\theta}^T \frac{1}{n} \tilde{\mathbf{X}}^T \tilde{\mathbf{X}} \boldsymbol{\theta} \leq \kappa_u$.

Condition 3. The error vector $\boldsymbol{\epsilon}_j$ has a mean of zero and sub-Gaussian tails for all $j \in \{1, \ldots, r\}$. That is, for all $j \in \{1, \ldots, r\}$ there exists a constant σ_j such that for any $\mathbf{a} \in \mathbb{R}^n$, with $||\mathbf{a}||_2 = 1$,

$$P(|\langle \boldsymbol{\epsilon}_j, \mathbf{a} \rangle| > t) \le 2 \exp\left(-\frac{t^2}{2\sigma_j^2}\right).$$

In addition, there exists a positive constant σ such that $\sigma \geq \max_{i} \sigma_{j}$.

Condition 4. There is at least one non-zero entry in β^* .

Conditions 1-3 are commonly made for penalized estimators, see Negahban et al. (2012) and the citations within. The framework in Negahban et al. (2012) does not directly apply to (7), because they assume the penalty is a norm which is not the case for our penalty, even for \mathbf{A} fixed, because of the inclusion of the Ridge penalty. However, under these conditions we can extend the approach of Negahban et al. (2012) and prove rates of convergence under

milder conditions then some similar methods (Price & Sherwood 2018, Li & Li 2010*b*). In our results the upper bound of the tuning parameter γ depends on the inverse of $||\beta^*||_{\infty}$, which is well defined under Condition 4.

Theorem 1. Assume Conditions 1-4 hold and $\gamma < \delta/(16r||\boldsymbol{\beta}^*||_{\infty})$ then for any $\mathbf{A} \in \mathcal{A}$

$$P\left[||\hat{\boldsymbol{\beta}}(\mathbf{A}) - \boldsymbol{\beta}^*||_2^2 \le \frac{9\delta^2}{4\kappa_l^2}s\right] \ge 1 - 2\exp\left[\frac{-n\delta^2}{32\sigma^2} + \log(rp)\right].$$

Proof of Theorem 1, and all other theoretical results, are provided in the supplemental material. Theorem 1 can be used to derive convergence rates for $\hat{\boldsymbol{\beta}}(\mathbf{A})$ for a set sequence of δ .

Corollary 1. Assume Conditions 1-3 hold, $\gamma < \delta/(16r||\boldsymbol{\beta}^*||_{\infty})$ and $\delta = \sqrt{\frac{64\sigma^2 \log(rp)}{n}}$ then for any $\mathbf{A} \in \mathcal{A}$

$$P\left[||\hat{\boldsymbol{\beta}}(\mathbf{A}) - \boldsymbol{\beta}^*||_2^2 \le 144 \frac{\sigma^2 \log(rp)}{n\kappa_l^2} s\right] \ge 1 - 2exp[-\log(rp)].$$

Corollary 1 provides that for fixed \mathbf{A} and $rp \to \infty$ that the estimator $\hat{\boldsymbol{\beta}}(\mathbf{A})$ achieves the same rate of converge as the lasso estimator (Negahban et al. 2012) and elastic net (Hebiri & van de Geer 2011). Using Corollary 1 and noting that $\hat{\boldsymbol{\beta}}$ is equivalent to a $\hat{\boldsymbol{\beta}}(\hat{\mathbf{A}})$ we can derive a rate of convergence for $\hat{\boldsymbol{\beta}}$, the global minimum of a nonconvex objective function.

Corollary 2. Assume Conditions 1-3 hold, $\gamma < \delta/(16r||\boldsymbol{\beta}^*||_{\infty})$ and $\delta = \sqrt{\frac{64\sigma^2 \log(rp)}{n}}$ then $P\left[||\hat{\boldsymbol{\beta}} - \boldsymbol{\beta}^*||_2^2 \le 144 \frac{\sigma^2 \log(rp)}{n\kappa_t^2}s\right] \ge 1 - 2exp[-\log(rp) + \log(|\boldsymbol{\mathcal{A}}|)].$

2.3 Model Selection Consistency

Let $\dot{\mathbf{X}}_{(1)} \in \mathbb{R}^{nr \times s}$ be the submatrix of $\tilde{\mathbf{X}}$ consisting of the active predictors of the regression defined in (6), and $\dot{\mathbf{X}}_{(2)} \in \mathbb{R}^{nr \times pr-s}$ be the submatrix of the remaining predictors. Define

 $\dot{\mathbf{X}} = (\dot{\mathbf{X}}_{(1)}, \dot{\mathbf{X}}_{(2)})$ and let $\dot{\boldsymbol{\beta}}^*$ and $\dot{\boldsymbol{\beta}}(\mathbf{A})$ be the similarly re-arranged versions of $\boldsymbol{\beta}^*$ and $\hat{\boldsymbol{\beta}}(\mathbf{A})$, respectively. In addition, let $\dot{\boldsymbol{\beta}}_{(1)}^*$ be the first *s* entries of $\dot{\boldsymbol{\beta}}^*$. Note, that we still have

$$\tilde{y} = \dot{\mathbf{X}}\dot{\boldsymbol{\beta}}^* + \tilde{\boldsymbol{\epsilon}} = \dot{\mathbf{X}}_{(1)}\dot{\boldsymbol{\beta}}_{(1)}^* + \tilde{\boldsymbol{\epsilon}}.$$

Thus model selection consistency results found for $\dot{\boldsymbol{\beta}}(\mathbf{A})$, also hold for $\hat{\boldsymbol{\beta}}(\mathbf{A})$.

Define $\dot{\mathbf{A}}$ such that it relates to $\dot{\mathbf{X}}$ the same way \mathbf{A} corresponds to \mathbf{X} . The signed graph Laplacian of $\dot{\mathbf{A}} \in \mathcal{A}$ is

$$\mathbf{L} = \dot{\mathbf{A}}^T \dot{\mathbf{A}} = \left(egin{array}{cc} \mathbf{L}_{11} & \mathbf{L}_{12} \ \mathbf{L}_{21} & \mathbf{L}_{22} \end{array}
ight),$$

where \mathbf{L}_{11} and \mathbf{L}_{22} are the signed graph Laplacians of the active and inactive sets respectfully. Finally, define $\mathbf{C}_{11} = \frac{1}{n} \dot{\mathbf{X}}_{(1)}^T \dot{\mathbf{X}}_{(1)}, \ \mathbf{C}_{22} = \frac{1}{n} \dot{\mathbf{X}}_{(2)}^T \dot{\mathbf{X}}_{(2)}, \ \text{and} \ \mathbf{C}_{21} = \mathbf{C}_{12}^T = \frac{1}{n} \dot{\mathbf{X}}_{(2)}^T \dot{\mathbf{X}}_{(1)}.$

Zhao & Yu (2006) first presented conditions for model selection consistency for lasso. While, Li & Li (2010*b*) present model selection results for a lasso-type estimator that includes a penalty for a graph structure in the predictors. The estimator in that paper is similar to the one in this paper, but with **A** fixed and a univariate response. The following conditions are generalizations of conditions presented in those papers. For vectors $\mathbf{a}, \mathbf{b} \in \mathbb{R}^d$, let sign(**a**) be a vector of the signs of the entries of **a**, |**a**| be the vector formed by taking the absolute values of each entry of **a**, min(**a**) is the smallest entry of **a** and **a** < **b** is true if and only if $a_i < b_i$ for all $i \in \{1, \ldots, d\}$.

Condition 5. Assume $\epsilon_i \sim N_r(\mathbf{0}_r, \sigma^2 \mathbf{D}_0)$, are *i.i.d* and $\mathbf{D}_0 \in \mathbb{R}^{r \times r}$ is a diagonal matrix where diagonal values are on the support (0, 1] and thus the maximum variance of the errors is σ^2 .

Condition 6. The minimum eigenvalue of C_{11} , k_1 , is positive for all n.

Condition 7. There exists a fixed η , that does not change with n, such that

$$\left| \left(\mathbf{C}_{21} + \frac{\gamma}{n} \mathbf{L}_{21} \right) \left(\mathbf{C}_{11} + \frac{\gamma}{n} \mathbf{L}_{11} \right)^{-1} \left\{ \operatorname{sign}[\dot{\boldsymbol{\beta}}_{(1)}^*] + \frac{\gamma}{\delta} \mathbf{L}_{11} \dot{\boldsymbol{\beta}}_{(1)}^* \right\} - \frac{\gamma}{\delta} \mathbf{L}_{21} \dot{\boldsymbol{\beta}}_{(1)}^* \right| \le \vec{1} - \eta.$$

Condition 7 is multivariate response version of the graph constrained irrepresentable condition from Li & Li (2010*b*), which is an extension of the original irrepresentable condition in Zhao & Yu (2006). Using these conditions, Lemma 1 from Li & Li (2010*a*) and additional conditions on the tuning parameters, we can prove the estimator is model selection consistent.

Theorem 2. Assume condition 1, 5, 6, and 7 hold. Define $\rho = \min \left[|(\mathbf{C}_{11} + \gamma/n\mathbf{L}_{11})^{-1}(\mathbf{C}_{11}\boldsymbol{\beta}_{(1)})| \right]$. If

$$\frac{\delta^2}{\log(rp-s)(n+\gamma^2 r/k_1)} \to \infty,$$

and

$$\frac{1}{\rho} \left\{ \sqrt{\frac{\log(s)}{nk_1}} + \delta \left\| (\mathbf{C}_{11} + \gamma/n\mathbf{L}_{11})^{-1} \operatorname{sign}[\dot{\boldsymbol{\beta}}_{(1)}^*] \right\|_{\infty} \right\} \to 0,$$

then for any \mathbf{A} ,

$$\lim_{n \to \infty} P\left\{ \operatorname{sign}[\dot{\boldsymbol{\beta}}(\mathbf{A})] = \operatorname{sign}(\dot{\boldsymbol{\beta}}^*) \right\} = 1.$$

Thus the estimator is is strongly sign consistent and therefore model selection consistency holds, see Zhao & Yu (2006) for a more detailed discussion of this relationship.

2.4 Post Selection Inference

This subsection provides details how to construct confidence intervals for non-zero coefficients of $\hat{\boldsymbol{\beta}}(\hat{\mathbf{A}})$ using the framework provided in Lee et al. (2016) for lasso estimators. Interpretation of models depends on the variables in the model and we restrict our analysis to inference conditional on the selected model. A consequence of this is $\boldsymbol{\beta}^*$ will change

depending on the model. Let $Q \subseteq \{1, \ldots, pr\}$ be a subset of selected predictor-response combinations. For simplicity of notation we assume that each response has at least one predictor selected. Define $\tilde{\mathbf{X}}_Q \in \mathbb{R}^{nr \times |Q|}$ to be the matrix made from the columns of $\tilde{\mathbf{X}}$ that belong to Q. Define $\boldsymbol{\mu} = E(\tilde{\mathbf{y}}) \in \mathbb{R}^{nr}$, $\tilde{\mathbf{X}}_Q^+ = \left(\tilde{\mathbf{X}}_Q^T \tilde{\mathbf{X}}_Q\right)^{-1} \tilde{\mathbf{X}}_Q^T \in \mathbb{R}^{|Q| \times nr}$ and

$$\boldsymbol{\beta}_{Q}^{*} = \underset{\boldsymbol{\beta} \in \mathbb{R}^{|Q|}}{\operatorname{arg min}} E \left\| \left\| \tilde{\mathbf{y}} - \tilde{\mathbf{X}}_{Q} \boldsymbol{\beta} \right\|_{2}^{2} = \tilde{\mathbf{X}}_{Q}^{+} \boldsymbol{\mu}.$$
(9)

Where $\boldsymbol{\beta}_Q^* = (\beta_{Q,1}^*, \dots, \beta_{Q,|Q|}^*)^T \in \mathbb{R}^{|Q|}$, is the true vector of the coefficients conditional on using only the covariates in Q. For purposes of post selection inference, we assume that $\tilde{\mathbf{y}}$ follows a normal distribution of $\tilde{\mathbf{y}} \sim N(\boldsymbol{\mu}, \boldsymbol{\Sigma}_{\boldsymbol{\epsilon}} \otimes \mathbf{I}_n)$.

Define the vector $\hat{\mathbf{s}} = (\hat{s}_1, \dots, \hat{s}_{pr})^T \in \mathbb{R}^{pr}$, where $\hat{s}_j = \operatorname{sign}(\hat{\beta}_j)$ if $\hat{\beta}_j \neq 0$ and $\hat{s}_j \in [-1, 1]$ if $\hat{\beta}_j = 0$. As defined in Tibshirani (2013) for the lasso estimator, for technical convenience we consider the equicorrelation set

$$\hat{Q} \equiv \{ j \in \{1, \dots, pr\} \mid |\hat{s}_j| = 1 \}.$$
(10)

The index of nonzero coefficients of $\hat{\beta}$ is a subset of \hat{Q} and typically these sets are the same.

Let \hat{Q} represent the selected model and let $\hat{\mathbf{A}} \in \mathbb{R}^{pr(r-1)/2 \times pr}$ be the graph selected from (7). Consider a specific index $j \in Q$, the goal of this work is to define an interval $C_q^{\mathbf{A},Q}$ for a given level α , such that

$$P(\beta_{Q,j}^* \in C_j^{\mathbf{A},Q} | \hat{Q} = Q, \hat{\mathbf{A}} = \mathbf{A}) \ge 1 - \alpha.$$
(11)

See Lee et al. (2016) for a justification of coverage of post-selection intervals being conditional on the selected model and a review of other approaches to post-selection inference. To derive this interval we consider an arbitrary vector of $\eta_Q \in \mathbb{R}^{nr}$ examine the conditional distribution of

$$\boldsymbol{\eta}_Q^T \tilde{\mathbf{y}} | (\hat{Q} = Q, \hat{\mathbf{A}} = \mathbf{A}).$$
(12)

Note that this definition does include all covariates that have non-zero coefficients, but could potentially include a non-zero coefficient as $\hat{s}_i \in [-1, 1]$. For $\hat{\beta}$ and \hat{s} to solve (8), conditional on $\hat{\mathbf{A}} = \mathbf{A}$, the following sufficient and necessary KKT conditions must be satisfied,

$$\left(\frac{1}{n}\tilde{\mathbf{X}}^{T}\tilde{\mathbf{X}} + \gamma \mathbf{A}^{T}\mathbf{A}\right)\hat{\boldsymbol{\beta}}(\mathbf{A}) - \frac{1}{n}\tilde{\mathbf{X}}^{T}\tilde{\mathbf{y}} + \delta\hat{s} = \mathbf{0}_{pr},\tag{13}$$

$$\hat{s}_j = \operatorname{sign}(\hat{\beta}_j) \text{ if } \hat{\beta}_j \neq 0,$$
(14)

and

$$\hat{s}_j \in [-1, 1] \text{ if } \hat{\beta}_j = 0.$$
 (15)

For a given \mathbf{A} , $\hat{\boldsymbol{\beta}}(\mathbf{A})$ is a unique minimizer of (8) and confidence intervals are constructed using these KKT conditions.

For the following results we need the OLS solution to be well defined and thus need $\tilde{\mathbf{X}}_Q^T \tilde{\mathbf{X}}_Q$ to be positive definite.

Condition 8. The matrix $\tilde{\mathbf{X}}_Q^T \tilde{\mathbf{X}}_Q$ is positive definite.

Define \mathbf{e}_j as the vector with zeros everywhere, but with a one in the *j*th position. The following theorem presents how to construct confidence intervals conditional on \hat{Q} and $\hat{\mathbf{A}}$.

Theorem 3. Let $F_{\mu,\sigma^2}^{[a,b]}(x)$ be the CDF of a random normal variable with mean μ , variance σ^2 , truncated to the interval [a,b]. Let $\boldsymbol{\eta} = \tilde{\mathbf{X}}(\tilde{\mathbf{X}}^T \tilde{\mathbf{X}})^{-1} \mathbf{e}_j$ and define L and U such that

$$F_{L,\boldsymbol{\eta}^{T}\tilde{\boldsymbol{\Sigma}}\boldsymbol{\eta}}^{[V_{\mathbf{s}_{Q},\mathbf{A}}^{-},V_{\mathbf{s}_{Q},\mathbf{A}}^{+}]}(\boldsymbol{\eta}^{T}\tilde{\mathbf{y}}) = 1 - \frac{\alpha}{2} \text{ and } F_{U,\boldsymbol{\eta}^{T}\tilde{\boldsymbol{\Sigma}}\boldsymbol{\eta}}^{[V_{\mathbf{s}_{Q},\mathbf{A}}^{-},V_{\mathbf{s}_{Q},\mathbf{A}}^{+}]}(\boldsymbol{\eta}^{T}\tilde{\mathbf{y}}) = \frac{\alpha}{2}$$

Where $V_{\mathbf{s}_Q,\mathbf{A}}^-$ and $V_{\mathbf{s}_Q,\mathbf{A}}^+$ are constants that depend on $\mathbf{s}_Q \in \mathbb{R}^{|Q|}$, \mathbf{A} , Q, $\tilde{\mathbf{\Sigma}}$, $\tilde{\mathbf{X}}_Q$ and \mathbf{y} . If Condition 8 holds then

$$P\left(\beta_{Q,j}^* \in [L,U] | \hat{Q} = Q, \hat{\mathbf{s}}_{\hat{Q}} = \mathbf{s}_Q, \hat{\mathbf{A}} = \mathbf{A}\right) = 1 - \alpha.$$

Theorem 3 provides a way to construct confidence intervals for a least squares model that accounts for the fact that variable selection was first done by satisfying the KKT conditions of the estimator $\hat{\beta}(\hat{\mathbf{A}})$. The result uses the framework proposed in Lee et al. (2016), which provides a more detailed discussion of the issue of post-selection inference and the intuition behind the interval construction. Theorem 3 assumes that $\tilde{\Sigma}$ is known, but usually in practice it will need to be estimated. If p < n, then the covariance matrix of the residuals from a saturated multivariate regression model could be used. An alternative approach would be to make some simplifying assumption to make it easier to estimate Σ , for instance assume it is diagonal.

2.5 Algorithm

To obtain the estimates for the estimator defined in (5) we propose using the formulation of the estimator that is similar to the optimization defined in (3). Define the following sets

$$P_{l} = \left\{ m : \|\boldsymbol{\beta}_{l} - \boldsymbol{\beta}_{m}\|_{2}^{2} \leq \|\boldsymbol{\beta}_{l} + \boldsymbol{\beta}_{m}\|_{2}^{2}, \|\boldsymbol{\beta}_{l} - \boldsymbol{\beta}_{m}\|_{2}^{2} < \|\boldsymbol{\beta}_{l}\|_{2}^{2}, m \in \{1, \dots, r\} \setminus \{l\} \right\}$$
$$N_{l} = \left\{ m : \|\boldsymbol{\beta}_{l} + \boldsymbol{\beta}_{m}\|_{2}^{2} < \|\boldsymbol{\beta}_{l} - \boldsymbol{\beta}_{m}\|_{2}^{2}, \|\boldsymbol{\beta}_{l} + \boldsymbol{\beta}_{m}\|_{2}^{2} < \|\boldsymbol{\beta}_{l}\|_{2}^{2}, m \in \{1, \dots, r\} \setminus \{l\} \right\}$$
$$Z_{l} = \left\{ m : \|\boldsymbol{\beta}_{l}\|_{2}^{2} \leq \|\boldsymbol{\beta}_{l} - \boldsymbol{\beta}_{m}\|_{2}^{2}, \|\boldsymbol{\beta}_{l}\|_{2}^{2} \leq \|\boldsymbol{\beta}_{l} + \boldsymbol{\beta}_{m}\|_{2}^{2}, m \in \{1, \dots, r\} \setminus \{l\} \right\}$$

such that $P_l \cup N_l \cup Z_l = \{1, \ldots, r\} \setminus \{l\}, P_l \cap N_l = \emptyset, P_l \cap Z_l = \emptyset$, and $Z_l \cap N_l = \emptyset$ $\forall l \in \{1, \ldots, r\}$. Solving (5) is equivalent to solving

$$\underset{\mathbf{B}\in\mathbb{R}^{p\times r},P_{1},N_{1},Z_{1},\dots,P_{r},N_{r},Z_{r}}{\operatorname{arg\,min}} \frac{1}{2n} \sum_{k=1}^{r} \sum_{i=1}^{n} (\mathbf{y}_{i} - \mathbf{x}_{i}^{T}\boldsymbol{\beta}_{k})^{T} (\mathbf{y}_{i} - \mathbf{x}_{i}^{T}\boldsymbol{\beta}_{k}) + \delta \sum_{k=1}^{r} \|\boldsymbol{\beta}_{j}\|_{1} + \frac{\gamma}{2} \sum_{k=1}^{r} \left(\sum_{l\in P_{k}} \|\boldsymbol{\beta}_{l} - \boldsymbol{\beta}_{k}\|_{2}^{2} + \sum_{m\in N_{k}} \|\boldsymbol{\beta}_{m} + \boldsymbol{\beta}_{k}\|_{2}^{2} + \sum_{s\in Z_{k}} \|\boldsymbol{\beta}_{k}\|_{2}^{2} \right).$$

$$(16)$$

By using this formulation we are able to propose an iterative algorithm that iterates between estimating the sets P_l, N_l , and Z_l , for each $l \in \{1, \ldots, r\}$, with the regression coefficients

fixed, and estimating the regression coefficients with the sets fixed. To initialize the algorithm estimates of **B** or P_l, N_l , and $Z_l, l = 1, ..., r$ are needed. We define initial values for the coefficients of the *c*th response are set by

$$\hat{\boldsymbol{\beta}}_{c}^{(0)} = \underset{\boldsymbol{\beta}_{c} \in \mathbb{R}^{p}}{\arg\min} \frac{1}{2n} \sum_{i=1}^{n} (y_{ic} - \mathbf{x}_{i}^{T} \boldsymbol{\beta}_{c})^{2} + \delta \|\boldsymbol{\beta}_{c}\|_{1} + \gamma \|\boldsymbol{\beta}_{c}\|_{2}^{2},$$
(17)

and define $\hat{\mathbf{B}}^{(w)} = \left(\hat{\boldsymbol{\beta}}_{1}^{(w)}, \dots, \hat{\boldsymbol{\beta}}_{r}^{(w)}\right)$ to be the *w*th iterative estimate of \mathbf{B}^{*} . Note we will also use $\hat{\boldsymbol{\beta}}^{(w)}$ to represent the vectorized version of $\hat{\mathbf{B}}^{(w)}$, consistent with previous notation. Given a fixed (δ, γ) we propose the following algorithm as a solution to (16).

- 1. Initialize $\hat{\boldsymbol{\beta}}_1^{(0)}, \dots, \hat{\boldsymbol{\beta}}_r^{(0)}$ as defined in (17).
- 2. For the wth iteration, where w > 0, repeat the following steps until the estimated sets P_l, N_l and Z_l do not change from iterate w - 1 to iterate w for all $l \in \{1, \ldots, r\}$.
 - (a) Holding $\hat{\mathbf{B}}^{(w-1)}$ fixed obtain estimates for $\hat{P}_k^{(w)}, \hat{N}_k^{(w)}, \hat{Z}_k^{(w)}$ for all $k \in \{1, \dots, r\}$ by solving the optimization

$$\underset{P_k, N_k, Z_k \forall k \in \{1, \dots, r\}}{\operatorname{minimize}} \sum_{k=1}^r \left(\sum_{l \in P_k} \| \hat{\boldsymbol{\beta}}_l^{(w-1)} - \hat{\boldsymbol{\beta}}_k^{(w-1)} \|_2^2 + \sum_{m \in N_k} \| \hat{\boldsymbol{\beta}}_m^{(w-1)} + \hat{\boldsymbol{\beta}}_k^{(w-1)} \|_2^2 + \sum_{s \in Z_k} \| \hat{\boldsymbol{\beta}}_k^{(w-1)} \|_2^2 \right).$$
(18)

(b) Holding $\hat{P}_k^{(w)}, \hat{N}_k^{(w)}, \hat{Z}_k^{(w)}$ for all $k \in \{1, \dots, r\}$ fixed obtain the estimate $\hat{\mathbf{B}}^{(w)}$ by solving the optimization

$$\underset{\mathbf{B}\in\mathbb{R}^{p\times r}}{\arg\min\frac{1}{2n}\sum_{k=1}^{r}\sum_{i=1}^{n}(\mathbf{y}_{i}-\mathbf{x}_{i}^{T}\boldsymbol{\beta}_{k})^{T}(\mathbf{y}_{i}-\mathbf{x}_{i}^{T}\boldsymbol{\beta}_{k})+\delta\sum_{k=1}^{r}\|\boldsymbol{\beta}_{j}\|_{1}} +\frac{\gamma}{2}\sum_{k=1}^{r}\left(\sum_{l\in\hat{P}_{k}^{(w)}}\|\boldsymbol{\beta}_{l}-\boldsymbol{\beta}_{k}\|_{2}^{2}+\sum_{m\in\hat{N}_{k}^{(w)}}\|\boldsymbol{\beta}_{m}+\boldsymbol{\beta}_{k}\|_{2}^{2}+\sum_{s\in\hat{Z}_{k}^{(w)}}\|\boldsymbol{\beta}_{k}\|_{2}^{2}\right).$$
(19)

The update steps, shown in (18) and (19) respectively, break the non-convex problem into two problems that can be solved directly with well studied solutions. The optimization in (18) can actually be solved directly in a single pass, where the set assignments to the w + 1 iteration for l = 1, ..., r are defined as

$$\begin{split} \hat{P}_{l}^{(w+1)} &= \left\{ m : \| \hat{\boldsymbol{\beta}}_{l}^{(w)} - \hat{\boldsymbol{\beta}}_{m}^{(w)} \|_{2}^{2} \le \| \hat{\boldsymbol{\beta}}_{l}^{(w)} + \hat{\boldsymbol{\beta}}_{m}^{(w)} \|_{2}^{2}, \| \hat{\boldsymbol{\beta}}_{l}^{(w)} - \hat{\boldsymbol{\beta}}_{m}^{(w)} \|_{2}^{2} < \| \hat{\boldsymbol{\beta}}_{l}^{(w)} \|_{2}^{2}, m \in \{1, \dots, r\} \setminus \{l\} \right\}, \\ \hat{N}_{l}^{(w+1)} &= \left\{ m : \| \hat{\boldsymbol{\beta}}_{l}^{(w)} + \hat{\boldsymbol{\beta}}_{m}^{(w)} \|_{2}^{2} < \| \hat{\boldsymbol{\beta}}_{l}^{(w)} - \hat{\boldsymbol{\beta}}_{m}^{(w)} \|_{2}^{2}, \| \hat{\boldsymbol{\beta}}_{l}^{(w)} + \hat{\boldsymbol{\beta}}_{m}^{(w)} \|_{2}^{2} < \| \hat{\boldsymbol{\beta}}_{l}^{(w)} \|_{2}^{2}, m \in \{1, \dots, r\} \setminus \{l\} \right\}, \\ \hat{Z}_{l}^{(w+1)} &= \left\{ m : \| \hat{\boldsymbol{\beta}}_{l}^{(w)} \|_{2}^{2} \le \| \hat{\boldsymbol{\beta}}_{l}^{(w)} - \hat{\boldsymbol{\beta}}_{m}^{(w)} \|_{2}^{2}, \| \hat{\boldsymbol{\beta}}_{l}^{(w)} \|_{2}^{2} \le \| \hat{\boldsymbol{\beta}}_{l}^{(w)} + \hat{\boldsymbol{\beta}}_{m}^{(w)} \|_{2}^{2}, m \in \{1, \dots, r\} \setminus \{l\} \right\}. \end{split}$$

Note that this update requires calculating r^2 terms and performing r(r-1) comparisons, but the sets are deterministic and no iterates are needed.

The optimization in (19) can be solved using a gradient descent algorithm where each β_{mk} is solved for iteratively with all other regression coefficients held fixed. Let $S(x, a) = \operatorname{sign}(x)(|x| - a)_+$ be the soft-thresholding operator. To solve (19) we use a coordinate descent algorithm where the update for the *m*th predictor for the *k*th response, $\bar{\beta}_{mk}$, is

$$\bar{\beta}_{mk} = \frac{S\left(n^{-1}\{\sum_{i=1}^{n} x_{im}y_{ik} - \sum_{j=1, j\neq m}^{p} (\sum_{i=1}^{n} x_{im}x_{ij})\bar{\beta}_{jk} + \gamma H_{mk}\}, \delta/2\right)}{n^{-1}\left(\sum_{i=1}^{n} x_{im}^{2} + \gamma\left\{(r-1) + \sum_{g=1}^{r} I\left[k \in \hat{P}_{g}^{(w)}\right] + \sum_{v=1}^{r} I\left[k \in \hat{N}_{v}^{(w)}\right]\right\}\right)}$$

where

$$H_{mk} = \sum_{l \in \hat{P}_{k}^{(w)}} \bar{\beta}_{ml} - \sum_{h \in \hat{N}_{k}^{(w)}} \bar{\beta}_{mh} + \sum_{g=1}^{r} \bar{\beta}_{mg} I\left[k \in \hat{P}_{g}^{(w)}\right] - \sum_{v=1}^{r} \bar{\beta}_{mv} I\left[k \in \hat{N}_{v}^{(w)}\right],$$

and $\bar{\beta}_{jv}$ are the current iterates of the coordinate descent algorithm. The algorithm iterates through all $m \in \{1, \ldots, p\}$ and $k \in \{1, \ldots, p\}$ in a similar fashion to other coordinate descent algorithms and converges with the change in the estimates across iterations is small. We propose selecting δ and γ using k-fold cross validation minimizing the validation

residual sum of squares. The proposed coordinate descent update is similar to GRACE, but differs due to structured use of the minimum penalty across multivariate responses.

While this algorithm is related to the iterative algorithms proposed by Witten et al. (2014), Price & Sherwood (2018), and Price, Molstad & Sherwood (2021), there are some differences. First, the penalties investigated by those authors are considered *cluster fusion penalties*. Second, the algorithms proposed in those papers are two stage procedure that require a solution to the well studied k-means problem, which can be unreliable and computationally burdensome in high-dimensional settings. The proposed algorithm provides an efficient algorithm that is guaranteed to find the optimal sets of the responses given the coefficients. An alternative approach would be an exhaustive search of all possible $\mathbf{A} \in \mathcal{A}$ which would provide a global minimizer. However, this becomes computational intractable for medium or large r as there are $3^{r(r-1)}$ combinations that must be searched before tuning parameter selection is considered. The iterative method we propose is not constrained in this way but is not guaranteed to achieve a global solution to the non-convex optimization problem. The proposed algorithm can be easily adjusted to accommodate if other penalty functions, besides the proposed ridge fusion, are used within the minimum function.

3 Extension to Multiple Binomial Responses

3.1 Method

This section details how to extend the proposed method to the setting where conditional on the predictors each response follows a binomial distribution. The focus will be on the binomial setting, but the method presented in this section can be generalized to multiple responses from the same exponential family with different mean functions. Define θ_k =

 $(\alpha_k, \boldsymbol{\omega}_k^T)$, where α_k is a response specific intercept, and $\mathbf{u}_i = (1, \mathbf{x}_i)$ be the p+1 dimensional vector of covariates for the *i*th observation. Let y_{ik} be a realization of the random variable

$$Y_{ik} \sim \operatorname{Bin}(n_{ik}, \pi_{ik}), \text{ where } \pi_{ik} = \frac{\exp(\mathbf{u}_i^T \boldsymbol{\theta}_k)}{1 + \exp(\mathbf{u}_i^T \boldsymbol{\theta}_k)}$$

We define the penalized likelihood for the minimum penalized model as

$$\sum_{k=1}^{r} \sum_{i=1}^{n} y_{ik} \mathbf{u}_{i}^{T} \boldsymbol{\theta}_{k} - n_{ik} \log \left[1 + \exp(\mathbf{u}_{i}^{T} \boldsymbol{\theta}_{k}) \right] + \delta \sum_{k=1}^{r} \|\boldsymbol{\omega}_{k}\|_{1} + \frac{\gamma}{2} \sum_{l=1}^{r} \sum_{m=1, l \neq m}^{r} \min \left(\|\boldsymbol{\theta}_{l} - \boldsymbol{\theta}_{m}\|_{2}^{2}, \|\boldsymbol{\theta}_{l} + \boldsymbol{\theta}_{m}\|_{2}^{2}, \|\boldsymbol{\theta}_{l}\|_{2}^{2} \right).$$

$$(20)$$

The lasso penalty does not include the intercept, because we assume that an intercept is part of the model for each response. However, the intercept is part of the penalty in the minimum function because we wish to group responses based on the relationships between fitted values.

3.2 Algorithm

We propose solving (20) by approximating it with a penalized quadratic function similar to the glmnet algorithm (Friedman et al. 2010). Define,

$$g(\pi_{ik}) = \log\left(\frac{\pi_{ik}}{1 - \pi_{ik}}\right) = \mathbf{u}_i^T \boldsymbol{\theta}_k.$$
 (21)

To implement our penalized quadratic approximation we define the following,

$$z_{ik} = g(y_{ik}) = g(\pi_{ik}) + \frac{y_{ik} - \pi_{ik}}{\pi_{ik}(1 - \pi_{ik})},$$
(22)

$$w_{ik} = \pi_{ik}(1 - \pi_{ik}),$$
 (23)

$$-l_k(\boldsymbol{\theta}_k) = \sum_{i=1}^n w_{ik} (z_{ik} - \mathbf{u}_i^T \boldsymbol{\theta}_k)^2.$$
(24)

Note that z_{ik} is the first order Taylor approximation of $g(y_{ik})$, and that w_{ik} is the conditional variance of z_{ik} given \mathbf{u}_i . Define $\mathbf{Z}_k = (z_{1k}, \ldots, z_{nk})^T \in \mathbb{R}^n$ and $\mathbf{W} = (w_{1k}, \ldots, w_{nk})^T \in \mathbb{R}^n$.

Let P_l, N_l and Z_l be defined as in Section 2.1, but now with respect to $\boldsymbol{\theta}_l$ instead of $\boldsymbol{\beta}_l$. We propose an iterative algorithm similar to that proposed in Section 2.5 to minimize (20).

Initial estimates of $\boldsymbol{\theta}_l$ or P_l, N_l , and Z_l are needed for all $l \in \{1, \ldots, r\}$. Similar to the method proposed in Section 2.5 we propose initializing the regression coefficients for each of the *r* responses separately using the elastic net estimator, which we define as $\hat{\boldsymbol{\theta}}_k^{(0)}$. Given a fixed (δ, γ) we propose the following algorithm as a solution to (24).

- 1. Initialize $\hat{\boldsymbol{\theta}}_1^{(0)}, \ldots, \hat{\boldsymbol{\theta}}_r^{(0)}$.
- 2. For the wth iteration where w > 0, repeat steps until the estimated sets P_l , N_l and Z_l do not change from iterate w 1 to iterate w:
 - (a) Holding $\hat{\boldsymbol{\theta}}_{k}^{(w-1)}$ fixed for all $k \in \{1, \dots, r\}$, obtain $\hat{P}_{l}^{(w)}$, $\hat{N}_{l}^{(w)}$ and $\hat{Z}_{l}^{(w)}$ for all $l \in \{1, \dots, r\}$ by solving the optimization $\sum_{k=1}^{r} \left(\sum_{k=1}^{r} \hat{\boldsymbol{\theta}}_{k}^{(w-1)} \hat{\boldsymbol{\theta}}_{k}^{(w-1)} \sum_{k=1}^{r} \hat{\boldsymbol{\theta}}_{k}^{(w-1)} \hat{\boldsymbol{\theta}}_{k}^{(w-1)}$

$$\underset{P_{1},N_{1},Z_{1},\dots,P_{r},N_{r},Z_{r}}{\text{minimize}} \sum_{k=1}^{r} \left(\sum_{l \in P_{k}} \|\hat{\boldsymbol{\theta}}_{l}^{(w-1)} - \hat{\boldsymbol{\theta}}_{k}^{(w-1)}\|_{2}^{2} + \sum_{m \in N_{k}} \|\hat{\boldsymbol{\theta}}_{m}^{(w-1)} + \hat{\boldsymbol{\theta}}_{k}^{(w-1)}\|_{2}^{2} + \sum_{s \in Z_{k}} \|\hat{\boldsymbol{\theta}}_{k}^{(w-1)}\|_{2}^{2} \right)$$
(25)

(b) Holding $\hat{P}_k^{(w)}, \hat{N}_k^{(w)}, \hat{Z}_k^{(w)}$ fixed for all $k \in \{1, \dots, r\}$ obtain the estimates of $\hat{\boldsymbol{\theta}}_k^{(w)}$ for all $k \in \{1, \dots, r\}$ by solving the optimization

$$\arg \min_{\boldsymbol{\theta}_{k} \in \mathbb{R}^{r}, \forall k \in \{1, \dots, r\}} \sum_{k=1}^{r} -l_{k}(\boldsymbol{\theta}_{k}) + \delta \sum_{k=1}^{r} \|\omega_{k}\|_{1} \\
+ \frac{\gamma}{2} \sum_{l=1}^{r} \left(\sum_{m \in \hat{P}_{l}^{(w)}} \|\boldsymbol{\theta}_{m} - \boldsymbol{\theta}_{l}\|_{2}^{2}, \sum_{h \in \hat{N}_{l}^{(w)}} \|\boldsymbol{\theta}_{h} + \boldsymbol{\theta}_{l}\|_{2}^{2}, \sum_{v \in Z_{l}^{(w)}} \|\boldsymbol{\theta}_{l}\|_{2}^{2} \right).$$
(26)

Similar to the algorithm in Section 2.5, this could be replaced with an exhaustive search algorithm for the global minimum. However, even for small r this will be computationally burdensome. The step for estimating the sets is identical to the previous algorithm and thus once again if the coefficients are considered given there is an easily derived global solution for the sets, unlike the K-means algorithm which has been used in similar problems.

For the update in (26) we propose using a proximal gradient descent method similar to the glmnet algorithm proposed by Friedman et al. (2008). Let \hat{w}_{ik} and \hat{z}_{ik} be the most recent estimates of w_{ik} and z_{ik} . The update of the *j*th variable of the *m*th response of the coefficients for the proposed proximal gradient descent method is

$$\bar{\theta}_{jm} = \frac{S\left(\sum_{i=1}^{n} \hat{w}_{ik} \{x_{ij}\hat{z}_{ik} - \sum_{l=1, l \neq j}^{p} u_{il}u_{ij}\bar{\theta}_{lm}\} + \gamma M_{jm}, I(j \neq 1)\delta/2\right)}{\left(\sum_{i=1}^{n} u_{ij}^{2} + \gamma \{(r-1) + \sum_{g=1}^{r} I(m \in \hat{P}_{g}^{(w)}) + \sum_{v=1}^{r} I(m \in \hat{N}_{v}^{(w)})\}\right)}$$

where

$$M_{jm} = \sum_{l \in \hat{P}_m^{(w)}} \bar{\theta}_{jl} - \sum_{h \in \hat{N}_m^{(w)}} \bar{\theta}_{jh} + \sum_{g=1}^r \bar{\theta}_{jg} I(m \in \hat{P}_g^{(w)}) - \sum_{v=1}^r \bar{\theta}_{jv} I(m \in \hat{N}_v^{(w)})$$

The algorithm iterates through all $j \in \{1, ..., p\}$ and $m \in \{1, ..., r\}$ until convergence and then updates the quadratic approximation and continues to solve the IRWLS optimization until convergence. This entire process is the solution to (26). To select δ and γ we propose using k-fold cross validation minimizing validation likelihood loss.

4 OLS Simulations

4.1 Data Generating Model and Evaluation Metrics

In this section we investigate the performance of the MinPen estimator defined by (5), the MinPen estimator with the correct indices matrix **A** known *a priori* (T-MinPen), the

multivariate cluster elastic net estimator (MCEN) Price & Sherwood (2018), the separate elastic net estimator (SEN) which fits the elastic net on each response using common tuning parameters, and joint elastic net estimator (JEN) which utilizes a group penalty on the same predictor variable across all responses (Friedman et al. 2008).

The SEN and JEN estimators are fit using the glmnet package in R. Tuning parameters are selected using a test train procedures, where each model is trained on n = 100observations and then evaluated on a different set of 100 test observations. The selected tuning parameters minimize the predicted sum of squares error on the test observations. We establish the T-MinPen estimator as a baseline for the MinPen estimator when the relationships would be known by a practitioner prior to fitting the model.

Let $\tilde{\Sigma}_X \in \mathbb{R}^{4 \times 4}$ with entries $\tilde{\sigma}_{ii} = 1$ and $\tilde{\sigma}_{ij} = \rho$, for $i \neq j$. The covariates are generated by $\mathbf{x}_i \sim N(\mathbf{0}_p, \mathbf{\Sigma}_x)$, where $\mathbf{\Sigma}_x$ is a block diagonal matrix with p/4 blocks of $\tilde{\mathbf{\Sigma}}_X$ with $\rho = 0.7$ on the diagonal blocks and all other entries are set to 0.

Define that data generating model to the be same as proposed in (1) such that $\epsilon_i \sim N_{15}(\mathbf{0}_{15}, \mathbf{I}_{15})$. In all simulations we perform 100 replications where each replication consists of using training data set with n = 100 and tuning parameters are selected using predicted squared error loss on 100 independently sampled observations. To evaluate the model we generate an independent validation set consisting of 1000 observations and calculate the average squared prediction error (SPE)

$$\frac{1}{15000} \sum_{k=1}^{15} \sum_{i=1}^{1000} (y_{ik} - \hat{y}_{ik})^2,$$

where y_{ik} is the kth element of \mathbf{y}_i and \hat{y}_{ik} is a prediction for the *i*th observation and kth response. We also use the mean square error (MSE) of the estimators of the regression coefficient matrix **B**, which we defined as

$$\frac{1}{15p} \sum_{k=1}^{15} \|\hat{\boldsymbol{\beta}}_k - \boldsymbol{\beta}_k^*\|_2^2$$

We also report on the number of true variables and false variables selected through true positive (TP) and false positive (FP) rates respectively over the 100 replications.

4.2 Block Structure in Regression Coefficients

In this simulation we will investigate how the methods perform when encountering a structured set of regression coefficients where

$$\mathbf{B}^{*} = \begin{pmatrix} \mathbf{\Delta}_{10}(\eta, \lambda) & \vec{0}_{10} & \vec{0}_{10} \\ \vec{0}_{10} & \mathbf{\Delta}_{10}(\eta, \lambda) & \vec{0}_{10} \\ \vec{0}_{10} & \vec{0}_{10} & \mathbf{\Delta}_{10}(\eta, \lambda) \\ \vec{0}_{p-30} & \vec{0}_{p-30} & \vec{0}_{p-30} \end{pmatrix}$$

We define the regression coefficients for subsets of responses as $\Delta_q(\eta, \lambda) = (-\eta_q - \lambda, \eta_q, \eta_q + \lambda, -\eta_q - 2\lambda, \eta_q + 3\lambda) \in \mathbb{R}^{q \times 5}$ where η_q is a q-dimensional vector with each element set equal to η and λ is a constant.

The proposed method should identify both the positive and negative relationships across the responses and leverage that information to improve prediction accuracy. Of the methods we compare to, only MCEN can identify relationships and it is limited to only identifying positive relationships. We investigate the settings where $(\eta, \lambda) \in$ $\{0.5, 1.0\} \times \{0.02, 0.05, 0.10\}$ for p = 40, 100, and 300. Figure 1 presents the results of the simulation for p = 300, while the results for p = 40 and 100 are shown in the supplemental material. The results show that MCEN and MinPen perform competitively in MSE and PSE for all values of η , while JEN is superior when $\gamma = 1$. With respect to true and

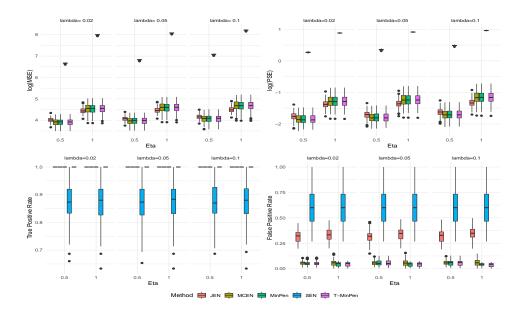


Figure 1: Results of the simulation described in Section 4.2 for the case of p = 300.

false positive rates, MinPen out performs or performs as well as the other approaches regardless of the values of η . It may be somewhat surprising that MCEN would perform well in this setting, but after investigation MCEN selects two clusters, a positive and negative cluster. As the effect size is consistent across each of these models, MCEN is competitive with MinPen though does tend to have a higher false positive rate.

The results presented show with respect to SPE all methods except for SEN perform similarly for all values of p, η , and λ . A similar results is found for both MSE and TP, with MinPen having a slight advantage for smaller p. The real differentiation between methods is shown when comparing the FP rate where we see MCEN and MinPen perform as well as T-MinPen and out perform all competitors SEN and JEN in the case of $\eta = 0.5$. In the case of $\eta = 1.0$ MinPen is able to perform closely to T-MinPen. For larger values of λ we see a much more consistent result in FP in MinPen than MCEN.

4.3 Overlapping Variables in Regression Coefficients

We next investigate a simulation where there is overlap between the non-zero variables in each coefficient vector. We define \mathbf{B}^* to be be structured such that the regression coefficients for the *k*th response, with r = 15, are defined as

$$\boldsymbol{\beta}_{k}^{*} = \begin{pmatrix} \vec{0}_{v(k-1)} \\ (-1)^{k} \vec{0.5}_{10} \\ \vec{0}_{p-v(k-1)-10} \end{pmatrix},$$

We investigate the settings $(v, p) \in \{0, 2, 4\} \times \{100, 300\}$ over 100 replications. Figure 2 presents the case of p = 300, while the results for p = 100 are available in the supplemental material. When v = 0 each predictor is either active or inactive for all 15 responses. In this setting JEN out performs competitors in all metrics for every p studied, because it uses a group lasso penalty for each predictor across the fifteen different responses. As the amount of overlap increases MCEN and MinPen methods perform the best with regard to MSE and PSE. Similar to the simulations presented in Section 4.2 T-MinPen and MinPen out preform competitors finding fewer false positive variables. Thus using MinPen is superior as the method is able to detect the underlying structure in the regression coefficients more often than competitors.

5 Binomial Simulations

Next we study the impact of the minimum penalty methodology in binomial logistic regression setting and compare it to SEN and MCEN which also have binomial logistic regression cases for the respective methods. We will also compare against T-MinPen which is a version of MinPen where we fix the true \mathbf{A} to be known by a practitioner *a priori*. We generate

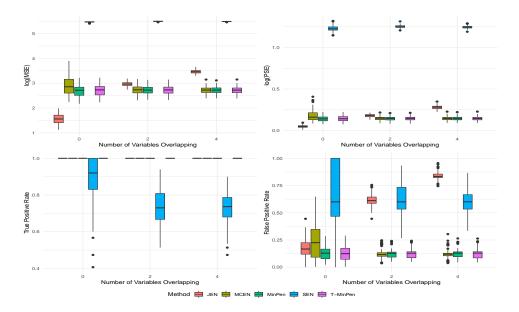


Figure 2: Results of the simulation described in Section 4.3 for the case of p = 300

 $\{\mathbf{x}_i\}_{i=1}^n$ in the exact same way as proposed in Section 4.1. The responses are generated by

$$y_{ik} \sim Bin(1, \pi_{ik}^*)$$
 where $\pi_{ik}^* = \frac{\exp(\mathbf{x}_i^T \boldsymbol{\beta}_k^*)}{1 + \exp(\mathbf{x}_i^T \boldsymbol{\beta}_k^*)}.$

In this simulation study we define $\beta_k^* k \in \{1, \ldots, r\}$ to be equivalent to the *k*th column of the **B**^{*} studied in section 4.2 for p = 40, 100, and 300 respectively. Again we study $(\eta, \lambda) \in \{0.5, 1.0\} \times \{0.02, 0.05, 0.10\}$. For each of the 100 iterations tuning parameters are selected using a train test procedure with 100 observations in the training set, and 100 observations in the test set. A validation of 1000 observations is to used evaluate methods using Kullback-Leibler divergence (KL), that is

$$\sum_{k=1}^{15} \sum_{i=1}^{1000} \left\{ \log \left(\frac{\hat{\pi}_{ik}}{\pi_{ik}^*} \right) \hat{\pi}_{ik} + \log \left(\frac{1 - \hat{\pi}_{ik}}{1 - \pi_{ik}^*} \right) (1 - \hat{\pi}_{ik}) \right\},\,$$

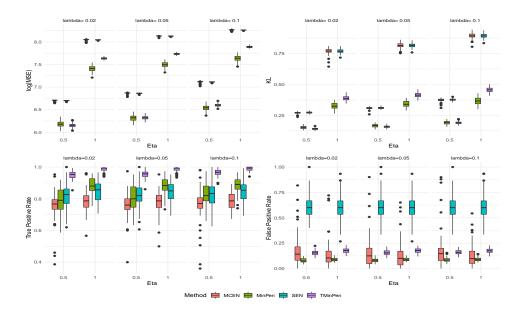


Figure 3: Results of the simulation described in Section 5 for p = 300.

where $\hat{\pi}_{ik}$ is the resulting estimated probability. We also compare methods using MSE, TP, and FP.

Figure 3 presents the results for the case of p = 300. The results for p = 40 and 100 are available in the supplementary material. The results show that MinPen performs as well as TMinPen and out performs the other competitors with respect to MSE and KL. We see that with regard to variable selection, MinPen is comparable, if not better, than the other methods with respect to TP and outperforms all approaches, even TMinPen, with respect to FP. TMinPen performs better the MinPen on TP but worse on FP indicating it is selecting more variables than necessary.

6 Applied Examples

6.1 Genomics Data

In this section we compare the post model selection inference approach outlined in this paper, after using the proposed least squares method, with the post model selection inference approach proposed by Lee et al. (2016), after using lasso for model selection. The models are fit to data analyzed by Votavova et al. (2011), who collected demographic, birth and gene expression data from 72 postpartum women and their newborns, but our analysis is limited to 64 of the women, 65 had complete data and one was dropped due to outliers in the demographic data. Four response variables are modeled: placental weight, newborn weight and two measures of cotinine level, one from the mother's peripheral blood and a second from the umbilical cord. The predictors are Smoking status, mother's age, mother's BMI, parity, gestational age and 33 gene expression probes. The 33 probes were selected by taking the absolute value of the correlation for each response and the 24,526 probes measured in the study and using a union of the top ten for each response.

The lasso models are fit separately for each response. For both methods tuning parameters are selected to minimize the mean squared prediction error from five folds cross validation. For post selection inference, $\tilde{\Sigma}$ is estimated using the covariance matrix of residuals from the full multivariate linear regression model. The four responses are related. First, the two cotinine measurements are measurements of the same variable at nearly the same time, but from different samples. In addition, cotinine levels are high in people who smoke and it has been shown that lower cotinine levels are associated with larger birth weights (Li et al. 1993). In addition, Wang et al. (2014) found both smoking has a negative relationship with both birth and placenta weights. Finally, newborn and placental weights are positively correlated (Molteni et al. 1978). The analysis of this data set serves two pur-

poses. First, we want to compare post selection inference results with an existing method. Second, we want to verify that minimizing (7) with real data can provide a sensible graph of the responses. For the latter issue the answer is yes. Table 1 provides the entries of $\hat{\mathbf{A}}$ and aligning with the scientific literature the optimal penalty is to penalize a difference in the weight measurements and a difference in the cotinine measurements, while all other comparisons the penalty enforces a negative correlation between the measurements.

	Placental Weight	Newborn Weight	Blood Cotinine	Cord Cotinine
Placental Weight	0	-1	1	1
Newborn Weight	-1	0	1	1
Blood Cotinine	1	1	0	-1
Cord Cotinine	1	1	-1	0

Table 1: The A matrix for the genomics data

Figure 4 provides the post selection confidence intervals for the lasso and MinPen methods. The selected models are very similar with the biggest model difference being in the model of the weight for placenta, where MinPen selects four more variables include the mother's age which is the only demographic variable difference between the models. In addition, for this model the post selection results for MinPen find a negative relationship between smoking and placenta weight, but the lasso results are inconclusive. The previously cited literature appears indicates that for this particular relationship the lasso approach suffers from a Type II error, while MinPen correctly identifies this relationship. However, for modeling cotinine level in the umbilical cord, the one setting where the variables selected are the same, the lasso confidence intervals are noticeably smaller.

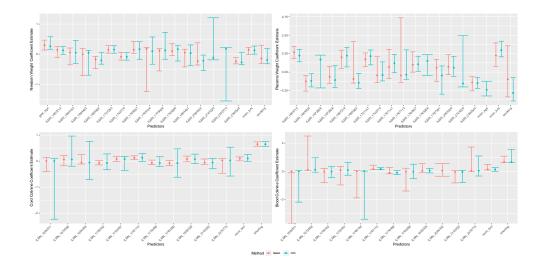


Figure 4: Post Selection Confidence Intervals: Clockwise starting in the top left corner; (1) Newborn weight; (2) Placenta Weight; (3) Umbilical cord cotinine level; (4) Peripheral blood cotinine level

6.2 Connecticut Multiple Substance Use Data

We investigate the use of the minimum penalty to identify co-occurring drug use based on accidental drug related deaths in Connecticut from 2012-2018. The data set consists of 5097 accidental overdoses in the state of Connecticut and corresponding toxicology, death certificate, and scene investigation information from the Chief Medical Examiner Office, publicly available at https://data.ct.gov and included in the supplemental material. We identify 17 response variables which are 16 individual drugs that were investigated plus a response that identifies if any opiate is involved. The goal of this analysis is to identify how different drugs appear (or do not appear) in cases of overdose, with a hope of better understanding and being able to predict co-occurrence of drugs. By being able to predict co-occurrence of drugs, both clinical practitioners and law enforcement are better able to

understand emerging trends in drug usage allowing them to quickly identify, intervene, and mitigate issues.

The covariates in this data are indicators of the city and state of the residence and death, along with using a bag of words approach to find common language used in the cause of death and description of injury provided by police and the medical examiner. As the goal is to better understand the relationships between drugs being used, information in the cause of death may not be relevant to all drugs found in the system. In total 111 covariates were used in this analysis. A validation set was created from 500 randomly selected observations from 2018 tuning parameters were selected using a 90/10 train/test approach using total classification error rate. Once the tuning parameters were obtained the model was fit using the chosen tuning parameters on all data less the validation set. We fit MinPen, MCEN, and elastic net models using a common sparsity parameter for all 17 responses, and compare methods using ROC curves on the validation set, see Figure 5. The results show the MinPen is competitive or out performs MCEN and the elastic net approach. Furthermore, MinPen provides relationships between responses. Only positive or no relationships were found between responses, that is no negative relationships were found. The supplementary material includes a graph of these relationships. Most notably heroin is not predictive or related to fentanyl, fentanyl analogue, benzodiazapine, or inform the presence of other opioids. The opioid variable was mostly related to prescription based drugs such as oxymorphone, hydrocodone, methadone, among others. Methadone was positively related to every other variable which is notable because it is used as a medical assisted treatment for opiate abuse. Further, the model finds benzodiazapine is not related to cocaine, which is of interest as benzodiasipine is used to treat cocaine toxicity. The results show MinPen provides better prediction accuracy than both MCEN and elastic net, providing evidence that use of the minimum penalty can be beneficial in the presence

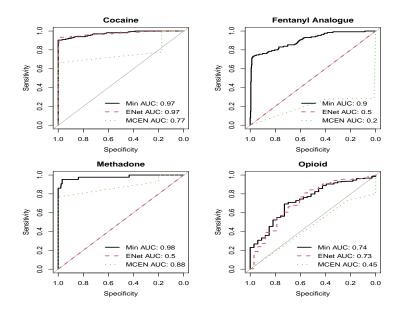


Figure 5: Results of the validation set of 500 observations from 2018 with regard to four substances: Cocaine, Fentanyl Analogue, Methadone, and the ability to detect any opioid. The results show that MinPen performs as well if not better with regard to AUC in classification.

of complex and potentially unknown relationships. In the supplemental material we also present a heatmap of the coefficients produced by MinPen for each response.

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