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# Combining $l^1$ and Higher Order $l^p$ Penalization in Regression Models

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#### Abstract

We create the package Penalized Regression on Steroids (**pros**) to combine  $l^1$  penalized regression with higher order  $l^p$  penalizations built upon the Elasticnet idea. The package is able to fit regression models with penalizations ranging from  $l^1$  to  $l^{10}$ .

Keywords: Penalized Regression, C++, Python, R.

## 1. Introduction

In statistics and probability theory it is common to impose moment assumptions on a random variable  $X:\Omega\to\mathbb{R}^n$  such as  $E(|X|^k)<\infty$  for  $k\in\mathbb{N}$ . These constraints correspond to the  $L^p$  spaces which allow control over the width and the height of such random variables. Constraints of this type may also be motivated geometrically. Consider for example an Elasticnet penalty  $Q(x)=\frac{1}{2}|x|+\frac{1}{2}|y|+\frac{1}{2}|x|^2+\frac{1}{2}|y|^2\leq 1$  shown on the left and a new penalty  $P(x)=\frac{1}{2}|x|+\frac{1}{2}|y|+\frac{1}{2}|x|^4+\frac{1}{2}|y|^4\leq 1$  shown on the right.

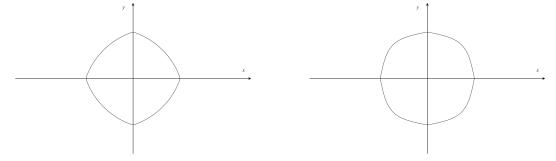


Figure 1: A particular ElasticNet penalty shown on the left and a new penalty shown on the right shown to "bow" out the penalization while retaining convexity.

A scientist may want the ability to control the height and width of the penalization while still retaining sparse solutions. In this project, we build the Penalized Regression on Steroids (**pros**) package available to the R programming language to expand upon the idea of ElasticNet by Zou and Hastie (2005), which we believe approximately encapsulates this idea.

## 2. Penalized Regression Implementation

In this section, we discuss how the package was implemented. In particular, the construction of the problem and the algorithms used. Let

$$L_{\lambda}(\beta) = \frac{1}{2} \|y - X\beta\|_{2}^{2} + \lambda P(\beta)$$

be the objective to minimize with penalty

$$\lambda P(\beta) = \lambda \alpha_0 \|\beta\|_1 + \lambda \sum_{k=1}^{5} \alpha_k \|\beta\|_{2k}^{2k}$$

where  $y \in \mathbb{R}^n$ ,  $X \in M_{n \times p}(\mathbb{R})$ , and  $\beta \in \mathbb{R}^p$ ,  $\lambda \in \mathbb{R}_+$  and  $\alpha$ 's are convex combinations. An important property of this penalty is convex and completely separable which allows for coordinate-wise optimization. In **glmnet** by Simon, Friedman, Hastie, and Tibshirani (2011), the authors use coordinate descent. Further, with the Elasticnet penalty by Zou and Hastie (2005), the coordinate wise minimization yields an analytic, closed form solution at each iteration avoiding any need for a line search or step size. The penalty that we propose lacks an analytic solution and thus a new algorithm is needed.

## 2.1. Optimization Algorithms and Implementation

The first algorithm utilized was the subgradient coordinate algorithm shown below.

## Algorithm 1: Subgradient Coordinate Algorithm

```
Choose \beta^0 \in \mathbb{R}^p, tolerance \delta > 0, R > 0, and maximum iterations N.

Set k \leftarrow 0

Set the step size h \leftarrow \frac{R}{\sqrt{1+N}}

repeat
\begin{array}{c|c} \text{Permute } I = \{1, \dots, p\} \\ \text{for } i \in I \text{ do} \\ \beta_i^{k+1} \leftarrow \beta_i^k - hg^i \text{ where } g_i^k \in (\partial L_{\lambda}(\beta^k))_i \\ k \leftarrow k + 1 \end{array}
```

**until** Until the objective difference is less than  $\delta$ ;

The drawbacks of this algorithm include lack of the descent property, no good stopping criterion, and many possible choices for the subgradient. The step size is optimal and chosen due to Nesterov (1998) with a worst case convergence rate of  $O(\frac{1}{\sqrt{k}})$ . Ultimately, no line search can be implemented and the step size must be tuned by the user, which may be difficult for many users. The stopping criterion is also expensive at  $O(n^2)$  flops at each iteration. Due to the separability of the penalization, a better algorithm is proximal gradient coordinate

Due to the separability of the penalization, a better algorithm is proximal gradient coordinate descent shown below.

## Algorithm 2: Proximal Gradient Coordinate Descent

```
Choose \beta^0 \in \mathbb{R}^p and tolerance \delta > 0;

Set k \leftarrow 0

repeat

Randomly permute I = \{1, \dots, p\}

for i \in I do

Set the step size h_i > 0 or use line search.

\beta_i^{k+1} \leftarrow (\mathbf{prox}_{h_iL})_i(\beta_i^k - h^k \langle X_i, y - X\beta \rangle)

k \leftarrow k + 1

end

until Until the Moreau-Yoshida mapping M_{h_k,f} < \delta;
```

The major benefit of this algorithm is that it is a descent method and thus line search may be implemented. Moreover, the stopping criterion is cheap to compute at O(p) flops. The Moreau-Yoshida map recovers the worst case convergence of  $O(\frac{1}{k})$  generally and coordinate descent algorithms are explored by Nesterov (2010).

## 2.2. Cross-Validation Algorithms and Implementation

In order to tune the Lagrangian dual variable  $\lambda$ , K-fold cross validation is implemented. To improve the speed of cross-validation, a warm start algorithm is used defined below.

## Algorithm 3: Warm Start Cross-Validation

```
Choose a sequence of Langrangian dual variables \lambda_1, \ldots, \lambda_N, and initial value \beta^0. Order \lambda_{(1)}, \ldots, \lambda_{(N)} descending. \beta^{Warm} \leftarrow \beta^0 for k \in 1, \ldots, N do \beta^k \leftarrow by Cross-Validation with \lambda_{(k)} warm started with \beta^{Warm}. \beta^{Warm} \leftarrow \beta^k end
```

A difficult problem in general is choosing the sequence of  $\lambda$ 's. In **glmnet**, the default is to choose a sequence of length 100 starting from the first  $\lambda_{max}$  where  $\beta$  is completely zero to the lower value  $.001\lambda_{max}$  on a log-scale. The lower  $\lambda$  value in the sequence is a real difficulty and this implementation does not fully address this. Due to time, we implement a similar default choice. We choose the lower  $\lambda = cp$  where c > 0 is a tuning parameter and then create a sequence of length 100 from this lower value. The default choice comparisons are explored in a subsequent section.

## 2.3. Step Size Rules and Line Search Implementation

The lack of an analytic solution introduces a necessary step size rule in order to fit a regression model with the new penalization. A good line search method due to Beck and Teboulle (2009) requires an evaluation of the objective function over the entire dataset at each iteration. This has a computational complexity of  $O(n^3)$  flops at each iteration. This was attempted, but never fully implemented in this package for this reason. Instead, a user defined step size rule is required. Thus, poor step size choices do not guarantee convergence. This is a suboptimal choice, but allows the potential for the package to be applied to very large datasets.

We introduce an alternative algorithm to solve this issue described below.

## Algorithm 4: Step Size Tuning Avoidance Algorithm

Choose initial value  $\beta^0$ , initial step size h, and maximum iterations N. repeat

```
h \leftarrow \frac{1}{2}h
 \beta \leftarrow by Proximal Gradient Descent with step size h
until Until Proximal Gradient Descent convergences;
```

If the maximum iterations for the algorithm are not too large, this may allow users to avoid having to tune a step size while retaining good performance. Currently, this is not implemented due to time constraints for the project.

## 3. The Penalized Regression on Steroids Package

## 3.1. Package Architecture

The package **pros** is built using C++ and the Eigen library by Guennebaud, Jacob *et al.* (2010) is utilized for fast matrix computations. This is analogous to using Fortran with LA-PACK by Anderson, Bai, Dongarra, Greenbaum, McKenney, Du Croz, Hammarling, Demmel, Bischof, and Sorensen (1990). The **pros** library is not dependent on R. To interact with R, 2 interfacing layers are needed: an R to C interface and an R program defining user callable functions. The benefit is that many other popular programming languages can be interfaced easily. The following diagram illustrates this idea showing that the **pros** package is more general than just an R package.

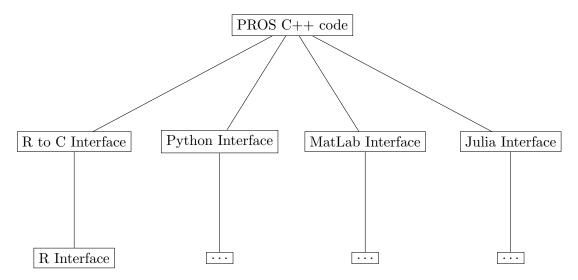


Figure 2: Illustrates interfacing the **pros** package with other languages.

The R interface in its simplest form is comprised of 2 functions. A single fitting function with prediction

```
> fit <- pros(X, y, alpha)</pre>
```

```
> predict(fit, X)
```

and a cross-validation function with prediction

```
> cv <- cv.pros(X, y, alpha)
> predict(cv, X)
```

We illustrate the usage of these functions in the following sections.

## 3.2. The Regression Fitting Function

The **pros** function is used to fit a single regression model with a specified penalization. The signature of this function at the time of this paper is as follows:

- X is an  $n \times m$ -dimensional matrix of the data.
- y is an *n*-dimensional vector of response values.
- alpha is a 6-dimensional vector of the convex combination corresponding to the penalization:
  - $-\alpha_1$  is the  $l^1$  penalty.
  - $-\alpha_2$  is the  $l^2$  penalty.
  - $-\alpha_3$  is the  $l^4$  penalty.
  - $-\alpha_4$  is the  $l^6$  penalty.
  - $-\alpha_5$  is the  $l^8$  penalty.
  - $-\alpha_6$  is the  $l^{10}$  penalty.
- lambda is the Lagrangian dual penalization parameter.
- step\_size is a tuning parameter defining the step size. Larger values are more aggressive and smaller values are less aggressive.
- algorithm is the optimization algorithm
  - proximal\_gradient\_cd uses proximal gradient coordinate descent.
  - subgradient\_cd uses subgradient coordinate descent.
- max\_iter is the maximum iterations the algorithm will run regardless of convergence.
- tolerance is the accuracy of the stopping criterion.
- random seed is the random seed used in the algorithms.

## 3.3. The Cross-Validation Function

The **cv.pros** function is used for K-fold cross-validation. The signature of this function at the time of this paper is as follows:

• X is an  $n \times m$ -dimensional matrix of the data.

- y is an n-dimensional vector of response values.
- alpha is a 6-dimensional vector of the convex combination corresponding to the penalization:
  - $-\alpha_1$  is the  $l^1$  penalty.
  - $-\alpha_2$  is the  $l^2$  penalty.
  - $-\alpha_3$  is the  $l^4$  penalty.
  - $-\alpha_4$  is the  $l^6$  penalty.
  - $-\alpha_5$  is the  $l^8$  penalty.
  - $-\alpha_6$  is the  $l^{10}$  penalty.
- K\_fold is the number of folds in cross-validation.
- lambdas is a vector of dual penalization values to be evaluated.
- step\_size is a tuning parameter defining the step size. Larger values are more aggressive and smaller values are less aggressive.
- algorithm is the optimization algorithm
  - proximal\_gradient\_cd uses proximal gradient coordinate descent.
  - subgradient\_cd uses subgradient coordinate descent.
- max\_iter is the maximum iterations the algorithm will run regardless of convergence.
- tolerance is the accuracy of the stopping criterion.
- random seed is the random seed used in the algorithms.

## 3.4. CRAN Issues

At the time of this paper, the package in its current form will not be accepted onto the Comprehensive R Archive Network (CRAN). There are some minor changes needed to comply with the CRAN policies and one larger issue. In order to get good performance out of C++, the compiler needs to be aggressive. It seems that R allows many options for Fortran, but some modern options for C++ possibly need to be added. These options need to be available in general so the author will reach out to the R developers about this.

## 4. Numerical Examples

## 4.1. The Analysis Procedure

The analysis in the following experiments is performed in the following way:

- The data is split into a training and test.
- The predictor data is standardized.

- The **glmnet** library and the **pros** library were used for fitting. The **glmnet** gives varying results for a single seed so we report a range of values.
- 10-fold cross-validation was used to tune the Lagrangian dual variable  $\lambda$ .
- The metric for comparison is the test mean squared error defined as

$$\frac{1}{n} \sum_{k=1}^{n} \left( y_i - \hat{\beta}_0 1 - X_i^T \hat{\beta} \right)^2$$

where  $\hat{\beta}_0$ ,  $\hat{\beta}$  are fit over the training set and y and X are the response and predictor matrix belonging to the test set.

• The source code for this analysis is openly available at the **pros** repository available here Brown (2018).

## 4.2. The Boston Housing Dataset

The Boston Housing dataset is popular from Harrison and Rubinfeld (1978). There are 13 predictors and the response is the median value of owner-occupied homes in \$1000's. The data is randomly split into a training set with 404 observations and a test set with 102 observations. We compare default implementations between the two libraries:

Penalty	Tuning	Test Mean Squared Error
Lasso (glmnet)	$\alpha = 1$	26.96866-28.45781
ElasticNet (glmnet)	$\alpha = 1/2$	26.55704-26.86488
Lasso (pros)	$\alpha = 1$	26.47315
ElasticNet (pros)	$\alpha = 1/2$	26.07919
4th Moment (pros)	$\alpha = 1/2$	25.5008
10th Moment (pros)	$\alpha = 1/2$	30.54759

Next, we compare manually tuned results using **pros**:

Penalty	Tuning	Test Mean Squared Error
ElasticNet (pros)	$\alpha = 0.1$	25.52578
4th Moment (pros)	$\alpha = 0.6$	25.44149
10th Moment (pros)	$\alpha = 0.9$	25.68279

## 4.3. The Prostate Dataset Analysis

This is a popular dataset from A. Stamey, N. Kabalin, E. Mcneal, M. Johnstone, Freiha, A. Redwine, and Yang (1989) and analyzed in the ElasticNet paper Zou and Hastie (2005). There are 8 predictors and the response is the log of the prostate specific antigen. We compare default implementations between the two libraries:

Penalty	Tuning	Test Mean Squared Error
Lasso (glmnet)	$\alpha = 1$	0.4443432 - 0.4646501
ElasticNet (glmnet)	$\alpha = 1/2$	0.4539684 - 0.4633808
Lasso (pros)	$\alpha = 1$	0.4457657
ElasticNet (pros)	$\alpha = 1/2$	0.4637193
4th Moment (pros)	$\alpha = 1/2$	0.4621351

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Penalty	$\operatorname{Tuning}$	Test Mean Squared Error
ElasticNet (pros)	$\alpha = .98$	0.4442702
4th Moment (pros)	$\alpha = 0.96$	0.4441759
10th Moment (pros)	$\alpha = 0.96$	0.4441322

Next, we compare manually tuned results using **pros**:

#### 4.4. Discussion

One source of confusion where the author spent an embarrassingly long amount of time was noticing that **glmnet** uses a transformation for their  $\lambda$  values and these do not correspond with the typical Lasso model. Thus, direct comparisons are extremely confusing. Further complicating comparisons is that **glmnet** provides varying results for fixed random seeds. The author should have spent more time carefully reviewing the **glmnet** paper by Simon et al. (2011). Due to this, the values in the corresponding slides for this project are incorrect.

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