## SML: Exercise 2

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2022-11-07

### Introduction

This report aims to find the best prediction for past cumulative grocery sales (in dollars) for Dominick's Finer Foods based on an Elastic Net model by Zou and Hastie (2005).

#### Data

The data set contains seven years of store-level data collected at Dominick's Finer Foods by the University of Chicago Booth School of Business. The data can be found at https://www.chicagobooth.edu/research/kilts/datasets/dominicks. The data set contains 50 variables, which stem from:

- 1. customer count files, which contain information about in-store traffic;
- 2. a demographic file, which contains store-specific demographic data;
- 3. number identification files, which contain product information.

Of the fifty variables, GROCERY\_sum is used as dependent variable. Furthermore, four categorical variables are dropped; STORE, CITY, ZIP and SPHINDX. The remaining variables are potential predictor variables.

### Method

To find the optimal set of predictor variables, and there corresponding weights, we use a regression method that penalizes the size of coefficients. The penalty is useful when predictors are collinear, or the number of predictors destabilizes estimation. This data set only consists of 77 observations for 50 variables, hence the number of predictors would destabilize estimation if not penalized.

Let  $P(\beta)$  denote a general penalty function. Then, the loss function of the regression equation becomes

$$L(\beta) = (\mathbf{y} - \mathbf{x}\beta)^T (\mathbf{y} - \mathbf{x}\beta) + \lambda P(\beta), \tag{1}$$

where  $\lambda$  is the hyperparameter that determines the strength of the penalty. When  $P(\beta) = \beta^2$ , the regression is called 'ridge' regression. Similarly, when  $P(\beta) = |\beta|$ , the regression is called 'LASSO' regression. Finally, any convex combination  $P(\beta) = \alpha |\beta| + (1 - \alpha)\beta^2$  of the 'ridge' and 'LASSO' penalty, where  $\alpha$  denotes the weight on the 'LASSO' penalty, is called 'elastic net' (compare Zou and Hastie 2005). We use the elastic net to find the optimal set of predictor variables, since it exploits the LASSO property of variable selection, and the ridge . . .

We use an MM-algorithm to estimate the elasting net. The MM-algorithm uses a majorizing function to find the coefficient vector  $\beta$  that minimizes the loss function specified in Equation 1. This is because the minimum is not obtained analytically. Let  $\epsilon$  denote the desired level of precision. The majorizing function is specified as

$$L_{MM}(\beta) = \frac{1}{2} \beta^{\mathrm{T}} (\frac{1}{n} \mathbf{X}^{\mathrm{T}} \mathbf{X} + \lambda (1 - \alpha) \mathbf{I} + \lambda \alpha \mathbf{D}) \beta - \frac{1}{n} \beta^{\mathrm{T}} \mathbf{X}^{\mathrm{T}} \mathbf{y} + c,$$

where **D** is a  $p \times p$  diagonal matrix with elements

$$d_{jj} = 1/\max\left(|\beta_j^0|, \epsilon\right).$$

Furthermore,

$$c = \frac{1}{2n} \mathbf{y}^{\mathrm{T}} \mathbf{y} + \frac{1}{2} \lambda \alpha \sum_{j=1}^{p} |\beta_{j}^{0}|.$$

To find the optimal penalty strength  $\lambda$  and the weight of the elastic net  $\alpha$ , K-fold cross-validation is used. K-fold cross-validation starts with splitting the data set into K folds. Subsequently, K iterations are ran as follows: for each iteration  $k \in \{1, \ldots, K\}$ , fold k is the test set, whereas the remaining (K-1) folds together form the training set. On the training set, the elastic net is estimated for all  $\lambda_i \in (10^{-6+i})_{i=1}^{10}$ . Moreover, for each  $\lambda_i$ , all corresponding  $\alpha \in \{0, 0.01, \ldots, 1\}$  are estimated, such that there are 1000 combinations of the hyperparameters  $\alpha$  and  $\lambda$ . All estimated elastic nets (i.e., all combinations of the hyperparameters), are used to predict the dependent variable in the kth fold. When this procedure has been completed for all K folds, we have obtained fitted values for every observations in the data set, for all elastic nets. Consequently, prediction errors can be computed.

To evaluate which elastic net performs best (that is, the elastic net that maintains the optimal combination of hyperparameters), the Root Mean Square Error (RMSE) is computed for each elastic net; the elastic net that has the lowest RMSE is deemed to be the best model. Let  $\hat{\mathbf{y}}_{\text{test}(k)} = \mathbf{X}_{\text{test}(k)} \hat{\beta}_{\text{train}(k)}$  denote the fitted values, and  $n_k$  the number of observations in fold k. The RMSE is computed as

$$RMSE = \sqrt{\frac{1}{K} \sum_{k=1}^{K} MSE_k},$$

where

$$MSE = \frac{1}{n_k} (\mathbf{y}_{test(k)} - \mathbf{\hat{y}}_{test(k)})^{\mathrm{T}} (\mathbf{y}_{test(k)} - \mathbf{\hat{y}}_{test(k)}).$$

To validate the results of our MM model, we compare our coefficient estimates with the ones of the established glmnet algorithm. We do not expect the coefficients to match exactly, e.g. due to different optimization algorithms. Hence, we need to measure their differences. We do so with the Absolute Error (AE)

$$AE_i = \left| \hat{\beta}_i^{GLM} - \hat{\beta}_i^{MM} \right|,$$

and the Absolute Percentage Error (APE)

$$APE_i = \left| \frac{\hat{\beta}_i^{GLM} - \hat{\beta}_i^{MM}}{\hat{\beta}_i^{GLM}} \right|.$$

### Results

where  $\hat{\beta}_i^{GLM}$  and  $\hat{\beta}_i^{MM}$  are the i-th coefficients of the glmnet and MM model, respectively.

Indeed, when applying the glmnet and MM implementations to the Dominick's Finer Food data set, we get different coeffciients. Figure 1 shows the mean-absolute-percentage differences (MAPE) of these two implementations for the different predictors. For most predictors, the APE is around to 1%.

One explanation for this difference could lie in the algorithms' speed of convergence. For a given data set the MM algorithm might not converge fast enough to provide the same estimates as the glmnet implementation. In fact, the cyclical coordinate decent used in glmnet is a very efficient algorithm according to Friedman, Hastie, and Tibshirani (2010).

We investigate this hypothesis by looking at the estimates' differences for lower convergence thresholds  $\epsilon$  of the MM algorithm. Figure 2 shows the zero correlation between the Mean Absolute Error and the parameter

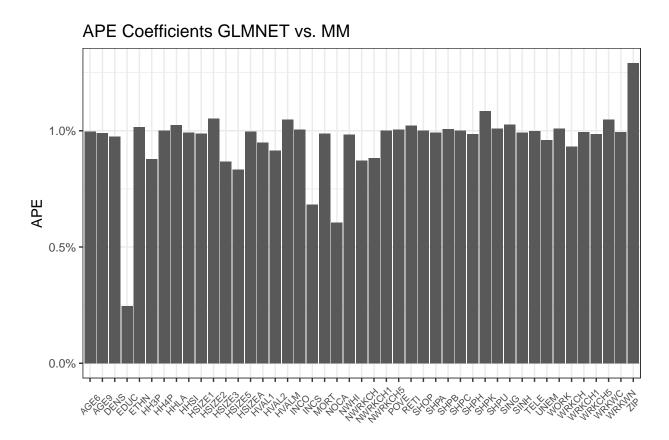


Figure 1: Absolute percentage difference between glmnet and MM coefficients.

# Median AE for decreasing $\boldsymbol{\epsilon}$ 1370917 -Щ 1370917 · 1370917 -1370917 -30 -70 -10 -20 -40 -50 -80 -90 0 -60 -100 $log_{10}(\epsilon)$

Figure 2: Development of AE fo decreasing precision threshold  $\epsilon.$ 

 $\epsilon$ : This precision parameter appears to play no role in the discrepancy between the coefficients of glmnet and the MM algorithm.

The uniform difference between the coefficients, and the result that this difference does not depend on the precision parameter  $\epsilon$  leads us to believe that there are some structural differences between the glmnet and MM algorithm. However, as these discrepancies are low and similar across predictors, both algorithms seem to implement the same model.

### Results

## Conclusion and Discussion

## References

## Code

Friedman, Jerome, Trevor Hastie, and Rob Tibshirani. 2010. "Regularization Paths for Generalized Linear Models via Coordinate Descent." *Journal of Statistical Software* 33 (1): 1.

Zou, Hui, and Trevor Hastie. 2005. "Regularization and Variable Selection via the Elastic Net." Journal of the Royal Statistical Society: Series B (Statistical Methodology) 67 (2): 301–20.