Linear Regresson: Regularization

Signal Data Science

Some helpful notes on the glmnet package are in a section at the end of this document. As you work through this assignment, you should **refer to those notes** to understand how glmnet() and cv.glmnet() work.

Exploring regularization with simulated data

Before using regularized linear regression on real data, we'll build some intuition by using regularization in a simpler context with simulated data. It's often useful to look at simple cases when trying to understand complex phenomena, so we'll consider the 1-dimensional case where we only have a single predictor variable.

Computational exercise

Define x and y using:

```
set.seed(1); j = 50; a = 0.25
x = rnorm(j)
error = sqrt(1 - a^2)*rnorm(j)
y = a*x + error
```

If you run summary $(lm(y \sim x - 1))$, corresponding to a linear model with no constant coefficient, you should get an estimated value of 0.2231 for a.

- Write a function cost(x, y, aEst, lambda, p) which takes
 - two vectors x and y of equal length,
 - a estimate of the value of a, aEst,
 - a regularization parameter lambda, and
 - a number p = 1 or 2, indicating whether L^1 or L^2 regularization is being performed.

Your function should return the sum of squared errors for the model y = aEst*x plus the L^p regularization term. Check that cost(1, 2, 3, 4, 2) returns 37.

- Create two vectors, one corresponding to values of λ given by $2^{-2}, 2^{-1}, \dots, 2^5, 2^6, 2^7$ and another corresponding to values of aEst from -0.1 to 0.3 in equally spaced increments of 0.001.
- Use expand.grid() on the two vectors of parameter to create a data frame called grid with two columns lambda and aEst, where each row is a unique pair of (lambda, aEst) values.
- Add costL1 and costL2 columns to grid where we'll store the cost of associated with each pair (lambda, aEst) for each of p = 1 and p = 2.
 Fill in those columns with cost().
- Write a function get_plot(lambda, p) which looks at the rows of grid with the specified value of lambda and returns the qplot() object generated by plotting the corresponding values of aEst on the abscissa (x-axis) and the corresponding values of either the L¹ or L² regularized cost function (depending on p) on the ordinate (y-axis).
- Use lapply() with get_plot() to create two lists, plotsL1 and plotsL2, where the *i*th plot is generated using the *i*th value of the vector of lambda values.
- Use multiplot with cols=2 and the plotlist parameter set to either plotsL1 or plotsL2 to visualize the results. Interpret the differences between the sets of plots for the L^1 and L^2 regularized cost function with respect to how L^1 regularization drives coefficient estimates to zero whereas L^2 regularization does not.

Theoretical explanation

We can see that L^1 regularization successfully drives coefficient estimates to 0 as λ increases while L^2 regularization does not. Why does this happen? We can get more insight into what's going on by looking at the underlying mathematics.

Suppose we have a vector of true values \mathbf{y} and a predictor variable \mathbf{x} , and consider an L^p regularized linear model for \mathbf{y} in terms of \mathbf{x} with regularization hyperparameter λ and coefficient estimate β . Call the sum of squared errors $SSE = S(\beta) = \sum_i (y_i - \beta x_i)^2$. Then our total cost function for the model is given by

$$C_n(\beta) = S(\beta) + \lambda |\beta|^p$$
.

First, let's consider the case when we perform L^2 regularization. In that situation, p = 2 so $|\beta|^p = \beta^2$, and our cost function is

$$C_2(\beta) = S(\beta) + \lambda \beta^2.$$

What value of β minimizes $C_2(\beta)$? Since $C_2(\beta)$ is the sum of two quadratic functions of β , it is smooth (being a quadratic function of β itself) and therefore the minimum is achieved when $C_2'(\beta) = 0$, *i.e.*, when

$$S'(\beta) + 2\lambda\beta = 0.$$

Remember that we're interested in the situation where regularization causes the coefficient estimate β to be driven to 0. It's then natural to ask: what needs to be true for $C_2(\beta)$ to be minimized at $\beta = 0$? The condition $C_2'(0) = 0$ must hold. Substituting $\beta = 0$ into our expression above, we obtain the condition

$$S'(0) + 2\lambda \cdot 0 = S'(0) = 0.$$

We can conclude that L^2 regularization will drive the coefficient estimate β to 0 *if and only if* the condition S'(0) = 0 holds. Since the sum of squared errors $S(\beta)$ is a smooth quadratic function of β , the condition S'(0) = 0 is equivalent to saying that the sum of squared errors is minimized at $\beta = 0$, *i.e.*, that \mathbf{y} is absolutely uncorrelated with \mathbf{x} .

Therefore: L^2 regularization drives the coefficient estimates to 0 if and only if the target variable is completely uncorrelated with its predictors. This is essentially *never* the case, so L^2 regularization will *never* drive coefficient estimates to 0.

Now, let's consider L^1 regularization, where p = 1 so

$$C_1(\beta) = S(\beta) + \lambda |\beta|.$$

This is the sum of a quadratic function of β and a scaled absolute value function of β . Each of the two functions has a single local minimum, so the global minimum of $C_1(\beta)$ must be located at *either* (1) at the minimum of $S(\beta)$, where $S'(\beta) = 0$, or at (2) the minimum of the regularization parameter, where $\beta = 0$.

Let's say that $S(\beta)$ is minimized at $\beta = \beta^*$. $C_1(\beta)$ will be minimized at the minimum of the regularization parameter, *i.e.*, at $\beta = 0$, if and only if $C_1(0) < C_1(\beta^*)$, *i.e.*,

$$S(0) < S(\beta^*) + \lambda |\beta^*|.$$

Rearranging terms, we can rewrite this inequality as

$$\lambda > \frac{S(0) - S(\beta^*)}{|\beta^*|}.$$

The right hand side of this inequality is a *constant* which depends solely upon \mathbf{y} and \mathbf{x} . As such, we can *always* make this inequality true by making the regularization hyperparameter λ sufficiently large!

Therefore: L^1 regularization can always drive a coefficient estimate to 0 by increasing the regularization hyperparameter λ past some constant threshold. In other words, with enough regularization we are *guaranteed* to be able to drive the coefficients of our model to 0.

Comparing regularization and stepwise regression

We will now begin to use the glmnet package, which provides the functions glmnet() and cv.glmnet() for regularized linear regression.

Continue using the aggregated speed dating dataset (speed-dating-simple) from yesterday. For simplicity, we'll restrict to analyzing average *attractiveness ratings* (attr_o) for *males*, making predictions for that specific rating in terms of the 17 self-rated activity variables.

Getting started with glmnet()

glmnet() can perform both L^1 and L^2 regularized linear regression as well as a mix of the two (which we'll be exploring later). When calling glmnet(), you can set alpha=1 for L^1 regularization and alpha=0 for L^2 regularization.

- Create a activities_scaled variable which is the result of calling scale() on a data frame containing only the 17 activity variables and an attr_o variable equal to the (unscaled) attr_o column of the original dataset. You'll be passing these variables into glmnet().
- Use glmnet() to fit both L^1 and L^2 regularized linear models for attroin terms of the variables in activities_scaled. By default, the function automatically determines a range of different λ values to try. To illustrate this, access the object returned by glmnet() and print out the values of λ used for each of the two regularized models.
- Write a function get_rmses(fit, df, target) that takes as input fit, the model object generated by a call to glmnet(), features, the predictors used in the call to glmnet(), and target, the true values of the target variable being predicted. Iterate over the values of λ which glmnet() tried. For each of those values, use predict() to generate predictions for the whole dataset and calculate the corresponding (non-cross-validated) RMSE. Return a vector of the calculated RMSE values.
- For both the L^1 and L^2 regularized linear fits, use get_rmses() to plot the RMSE corresponding to each value of λ against λ itself.

Making cross-validated RMSE estimates

We can see that the non-cross-validated RMSE is minimized at lambda = 0! This is not surprising, because *on the whole dataset* the linear fit which minimizes the RMSE is precisely (and almost tautologically) the one obtained by minimizing the sum of squared errors, without adding on any regularization term. However, the same cannot be said for the *cross-validated* RMSE estimates. Typically, adding in some regularization will reduce the amount of overfitting sufficiently well that the optimal value of λ is greater than 0.

We can automatically generate cross-validated error estimates for a range of different λ values with cv.glmnet(). For each value of λ tested, cv.glmnet() uses n-fold cross-validation (n=10 by default) to calculate an error estimate. After running fit = cv.glmnet(...), the value of λ (out of those tested) corresponding to the lowest error estimate can be accessed with fit\$lambda.min. Similarly, the entire range of λ values tested can be accessed with fit\$lambda and the cross-validated error estimates can be accessed with fit\$cvm. (See the notes on glmnet below for additional clarification.)

• Use cv.glmnet() to fit L^1 and L^2 regularized linear models for attractiveness ratings in terms of the 17 activity variables. For each one, plot the cross-validated error estimates against the values of λ tested. Interpret the results.

Stepwise regression vs. regularization

Stepwise regression tends to *overfit* because of problems with multiple hypothesis testing. It's easy to understand as an introduction to model comparison and serve as a good stepping stone to regularization, which is why we include it in our curriculum. However, *in practice*, regularizing a linear model is typically strictly superior to selecting a subset of the predictors with a stepwise method.

Read the entirety of this section before writing any code.

• Implement **20-fold** cross-validation. Within each fold, you should fit a model predicting attractiveness in terms of the 17 activities (1) with step() for backward stepwise regression and (2) with cv.glmnet() for both L^1 and L^2 regularized linear regression. Use the fitted model to make predictions on the held-out fold for all three methods. At the end of the 10 iterations, calculate and return the RMSE corresponding to each method. Interpret the results.

Keep the following in mind while you work:

¹Technically, by default cv.glmnet() calculates the *mean squared error* for linear regression models. However, since the square root function is monotonically increasing, it doesn't matter if one compares models using MSE or RMSE.

- After fitting a model with cv.glmnet(), be sure to pass s=fit\$lambda.min
 to predict(), because by default predict() does not use the value of λ
 which minimizes the cross-validated error.
- Within each cross-validation fold, you'll want to scale() the features which you pass into cv.glmnet(). However, when making predictions on the *held-out* data, you don't want to simply call scale() on that data directly; instead, you want to apply the *same transformations* which were applied to the training set. To do so, (1) extract the scaling parameters from the scaled training data (stored as attributes) and (2) pass them in as additional parameters to scale() when calling it on the test set.

Elastic net regression

Instead of penalizing the sum of squared errors by the L^1 or L^2 norm of the regression coefficients, we can penalize with a *combination* of the two, corresponding to setting the alpha parameter in glmnet() to a value *between* 0 and $1.^2$ This is known as elastic net regularization and usually performs better than pure L^1 or L^2 regularization alone. However, instead of simply finding a *single* hyperparameter λ , we now must find the optimal *pair* of hyperparameters (α , λ) (by testing a large number of such pairs and calculating the cross-validated RMSE corresponding to each one). This is a more difficult task, because we must search over a 2-dimensional space of hyperparameter combinations instead of a 1-dimensional space for the value of a single hyperparameter.

The caret package allows us to easily obtain a cross-validated estimate of the optimal (α, λ) values. Here's an example of how to use its train() function:

 $^{^2}$ Read the official documentation for glmnet to figure out how the α parameter works..

View the cross-validated RMSE estimates

caret_fit\$results\$RMSE

In the above example, we perform 10-fold cross-validation for each pair of hyperparameters (α, λ) to estimate the corresponding RMSE. The 10-fold cross validation is *repeated* three times, each time using a different random set of folds, in order to combat potential bias resulting from any particular choice of random folds. The optimal pair of values (α, λ) is the one corresponding to the lowest cross-validated RMSE.

• Use the caret package, following the above example, to find the optimal values for (α, λ) when predicting attractiveness ratings with elastic net regularization. Extract the minimum RMSE value obtained from the resulting caret_fit object and compare it to the cross-validated RMSE estimates obtained earlier with backward stepwise regression, L^1 regularized linear regression, and L^2 regularized linear regression.

A note on glmnet

Here, I'll cover two important points about the behavior of the glmnet package.

Passing in data

For lm(), you passed in the entire data frame, including both target variable and predictors. glmnet(features, target, ...) and cv.glmnet(features, target, ...) expect a *scaled matrix of predictors* for features and a numeric vector for target. Since scale() returns a matrix, you can just call scale() on a data frame of predictors and pass that in as features.

Picking values of λ

Ordinarily, one might expect that, for every different value of λ we want to try using with regularized linear regression, we would have to recompute the entire model from scratch. However, the glmnet package, through which we'll be using regularized linear regression, will automatically compute the regression coefficients for *a wide range of λ values simultaneously.³

When you call glmnet() – or, later, cv.glmnet() – you'll get out an object, which we'll call fit. (You should generally not be specifying which λ values the algorithm should use at this point – it'll try to determine that on its own.) By

³"Due to highly efficient updates and techniques such as warm starts and active-set convergence, our algorithms can compute the solution path very fast."

printing out fit in the console, you can see which values of λ were used by glmnet.

When you want to make predictions with this fit object, you'll have to specify which value of λ to use – instead of calling predict(fit, new_data), you'll want to call predict(fit, new_data, s=lambda) for some particular λ = lambda. Similarly, when extracting coefficients, you'll want to call coef(fit, s=lambda).

Finally, cv.glmnet() will use cross-validation to determine fit\$lambda.min and fit\$lambda.1se. The former is the value of λ (out of all those the algorithm evaluated) which minimizes the cross-validated mean squared error (MSE), and the latter is the greatest value of λ (again, of those evaluated by glmnet) such that the MSE corresponding to fit\$lambda.1se is within 1 standard error of the MSE corresponding to fit\$lambda.min.

If it turns out that the optimal value of λ lies at either end of the range of λ values used by glmnet, then you'll want to modify the range of λ . However, the documentation advises against passing in just a single value for the lambda parameter of glmnet() and cv.glmnet(), instead suggesting modifying nlambda and lambda.min.ratio.⁴ Nevertheless, there are times when passing in a single value makes sense, like when you've previously determined the optimal λ and want to just use that instead of a range of different λ values.

^{4&}quot;Typical usage is to have the program compute its own lambda sequence based on nlambda and lambda.min.ratio. Supplying a value of lambda overrides this. WARNING: use with care. Do not supply a single value for lambda (for predictions after CV use predict() instead). Supply instead a decreasing sequence of lambda values. glmnet relies on its warms starts for speed, and it's often faster to fit a whole path than compute a single fit."