Case 1

02582 Computational Data Analysis

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Introduction/Data description

We are given a set of data consisting of 100 observations (Y, X), where Y is the response (univariate) and X are the features (100-dimensional). We are additionally given a 1000 observations of the features.

The problem is then to build a predictive model, so that we can predict a new value of Y based on the 100 features X.

Y is a continuous variable, as are the feature variables X1 up to X95. X96 up to X100 are categorical variables, holding values of A, B, C and so forth. Since we do not know the real meaning behind those categorical values, we cannot assume that their order has any meaning, so they must be thought of as nominal.

Model and method

Since we are predicting a single continuous variable, a regression model seems like it would give us a good predictor. The approach is then to try out different regression methods, namely linear regression, ridge regression, lasso regression and elasticNet regression, and see which one can give us the best predictor for our data.

In short we load the data, handle missing values (see how in section below), one-hot encode the categorical data, standardize and normalize the data and then do K-fold cross-validation (K=10) for each of the regression methods. For each regression method, the cross-validation tries out different hyper parameters and picks the one that gives the model that has the best score (we decided on using the coefficient of determination, R^2 , as a scoring metric). We now have the best model we can find for each regression method, and out of these models we choose the model we think best fits our data (see how in section below).

Missing values

Missing numerical values are replaced by the mean of their variable (so for example a missing value in X1 is replaced by the mean of all the values in X1). Missing categorical values are replaced by the most common value of their variable (so for example, missing values in X96 are replaced by the most common value in X96, which is B). These are common practices which ensure that replacing missing values doesn't alter the underlying structure of the data set.

Factor handling

Factor variables, also called categorical variables, are handled by using one-hot encoding. One-hot encoding is a way to work with categorical data when doing regression. One-hot encoding transforms each categorical variable into multiple variables, 1 variable for each categorical value, and holding values of either 0 or 1 (1 if the categorical variable had said value in a particular row, 0 otherwise). One-hot encoding is illustrated in the figure below:

| X96 | X97 | X98 | X99 | X100 | one-hot | X96_A | X96_B | X97_A | X97_B | X97_C | |
|-----|-----|-----|-----|------|----------|-------|-------|-------|-------|-------|--|
| В | Α | С | С | Н | ====>>>> | 0 | 1 | 1 | 0 | 0 | |
| В | В | D | В | В | | 0 | 1 | 0 | 1 | 0 | |
| В | В | В | D | Ε | | 0 | 1 | 0 | 1 | 0 | |
| В | В | Α | В | D | | 0 | 1 | 0 | 1 | 0 | |
| Α | С | D | D | D | | 1 | 0 | 0 | 0 | 1 | |
| | | | | | | | | | | | |
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Figure 1: One-hot encoding explained

Model selection

Model selection is performed using cross-validated grid-search over a parameter grid. (https://scikit-learn.org/stable/modules/generated/sklearn.model_selection.GridSearchCV.html).

Model validation

Model validation is done by evaluating the coefficient of determination R^2 for each of the selected models for each regression method. R^2 is an evaluation metric that represents the ratio between how good a model is versus a model that always predicts the expected value of the dependent variable, disregarding the input features. The model with the highest R^2 (best possible score is 1.0) will be the best predictor for our data.

Results

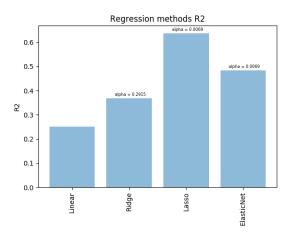


Figure 2: R^2 for the different regression methods tried

As can be seen by Figure 2, Lasso regression with hyperparameter α value 0.0069 yields the highest value for R^2 and is therefore the best and our chosen model. We will use this model to generate our predictions, \hat{y}_{new} .

To estimate RMSE, we use K-fold cross-validation again (K=10), but now on the 1000 extra observations and the predictions we just generated. For each of the K folds, we calculate the mean squared error, giving us an estimate of how good our predicted values, \hat{y}_{new} , are. We average over the K errors and take the square root of that average, obtaining an estimate of the *root mean squared error*, \widehat{RMSE} . We get that

 $\widehat{RMSE} = 0.38654208278816965$