Pipeline Optimization

Introduction:

In this assignment, I am going to optimize several ML models using Pipeline Optimization integrated with Bayesian optimization to find the best and optimum model for wine quality dataset. I am going to select several ML algorithms, apply the pipeline optimization on them, optimize them for their hyperparameters, and use them to predict the results. This configuration is supposed to be highly complicated. By comparing the results, we will find out which ML model shows better performance as well as providing insight for pipeline optimization process. Here, I used the regression learners including k-Nearest Neighbors (hyperparameters: k and distance), Random Forest (hyperparameters: num.trees, min.node.size, and mtry), Support Vector Machine (hyperparameters: cost, epsilon, and gamma), and Gradient Boosting (hyperparameters: nrounds and max_depth). Also, I applied scaling which scales features and factor encoding to each one of the learners. Root Mean Square Error was used as objective function and calculated for each iteration to find the model with minimum RMSE.

Dataset Description:

I am using the data from white wine which has 12 features and 4898 observations. Features include fixed acidity, volatile acidity, citric acid, residual sugar, chlorides, free sulfur dioxide, total sulfur dioxide, density, pH, sulphates, alcohol, and quality. I used all of the mentioned attributes to predict wine quality which has a numeric scale between 0 to 10. The dataset has no missing value. A summary of the dataset is presented in table 1.

Table 1. statistical summary of the white wine dataset

| Feature | Min | Mean | Max | SD |
|----------------------|------|-------|-------|-------|
| Fixed acidity | 3.80 | 6.85 | 14.20 | 0.84 |
| Volatile acidity | 0.08 | 0.27 | 1.10 | 0.10 |
| Citric acid | 0.00 | 0.33 | 1.66 | 0.12 |
| Residual sugar | 0.60 | 6.39 | 65.80 | 5.07 |
| Chlorides | 0.01 | 0.04 | 0.34 | 0.02 |
| Free sulfur dioxide | 2.00 | 35.31 | 289.0 | 17.00 |
| Total sulfur dioxide | 9.00 | 138.4 | 440.0 | 42.50 |
| Density | 0.98 | 0.99 | 1.04 | 0.003 |
| рН | 2.72 | 3.19 | 3.82 | 0.15 |
| Sulphates | 0.22 | 0.49 | 1.08 | 0.11 |
| Alcohol | 8.00 | 10.51 | 14.20 | 1.23 |
| Quality | 3 | 5.87 | 9 | 0.88 |

Experimental Setup:

I used R programming language and the libraries such as mlr3 (to implement ML tasks), mlr3pipelines (to set up the pipeline), mlr3learners (to train the models), mlr3tuning (to tune the hyperparameters), mlr3mbo (to implement Bayesian Optimization), mlr3viz (to visualize the results), readr (to read the data), caret (to split the data), and dplyr (to manage the data). I decided to fit the regression models to predict the wine quality, therefore, regression learners including k-Nearest Neighbors ("regr.kknn"), Random Forest ("regr.ranger"), Support Vector Machine ("regr.svm"), and Gradient Boosting ("regr.xgboost") were used in this exercise. I applied the scaling and factor encoding to each one of learners and made a graph for them (Figure 1).

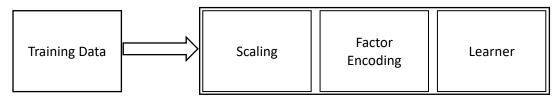


Figure 1. Schematic structure of pipeline optimization

The hyperparameters, related value ranges, and optimum value (results after optimization) for each ML model are presented in table 2. The Bayesian optimization approach was used in combination with pipeline optimization to minimize the objective function (which is RMSE).

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|---------------------------|----------------------------|----------|---------------|
| Model | Hyperparameter | Range | Optimum value |
| k-Nearest Neighbors | k | 1-20 | 12 |
| | distance | 1-20 | 4.85 |
| Random Forest | num.trees | 100-1000 | 315 |
| | min.node.size | 1-20 | 1 |
| | mtry | 1-10 | 2 |
| Support Vector Machine | cost | 0.1-10 | 1.99 |
| | epsilon | 0.01-2 | 0.23 |
| | gamma | 0.01-2 | 0.54 |
| Gradient Boosting | nrounds | 10-300 | 66 |
| | max_depth | 1-20 | 10 |

Table 2. Hyperparameters and related value range and best value for ML models

To tackle this problem and select the best ML algorithm after optimization process, I used the nested resampling by dividing the data into two sets (training and validation). I applied 5-fold cross validation to the training dataset and ran the experiment with 100 iterations for each model. At each iteration, the RMSE value was calculated and the settings with the lower RMSE value were selected. Afterward, I predicted the result for the validation set using optimized setting to

calculate the RMSE. Thus, we have RMSE for optimized model (training using 5-fold CV) and RMSE for validation to compare the models together, considering that the validation set prevents bias and overfitting.

Results:

The results of the optimization for training dataset (5-fold CV) were presented in figure 2. This plot illustrated the RMSE for each iteration and model. We can conclude that the Random Forest model is showing better performance and Gradient Boosting is the second best based on this figure. In addition, this figure demonstrates more fluctuation in comparison to the plot of hyperparameter optimization.

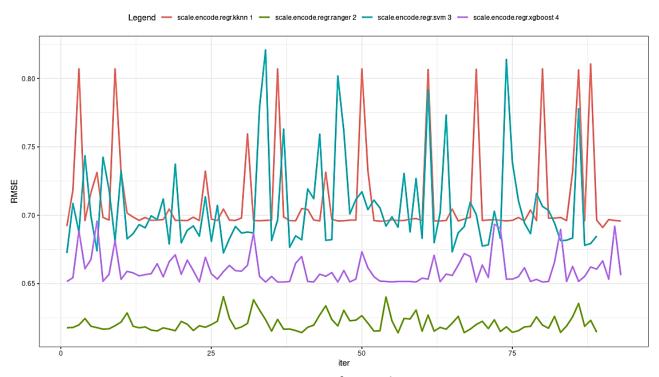


Figure 2. Optimization process for pipeline optimization.

The results for the best run after optimization for training (5-fold CV) and validations using optimum setting were presented in table 3.

| Model | RMSE for 5-fold CV | RMSE for validation |
|--------------------------|--------------------|---------------------|
| k-Nearest Neighbors | 0.69 | 0.70 |
| Random Forest | 0.61 | 0.60 |
| Support Vector Machine | 0.67 | 0.66 |
| Gradient Boosting | 0.65 | 0.64 |

We know the lower RMSE represents the better fit. Based on table 3, Random Forest shows better performance with RMSE = 0.60 in comparison with the other models. The RMSE of 5-fold CV and validation were approximately equal for all the models. All in all, we cannot observe any improvement in the results after combining pipeline and hyperparameter optimizations compared to previous exercise (hyperparameter optimization). It is suggested to repeat the experiment with specific pipeline for each ML algorithm.