Tutorial 10 - Cross-Validation & Evaluation

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Tutorial 10

This tutorial will cover out-of-sample predictions, cross-validatoin and evaluation metrics. You will learn:

- how to split your data into a test & training set
- how to train your model on the training data & make predictions on the test data
- how to calculate the out-of-sample RSME
- how to apply a cross-validation procedure
- how to get a confusion matrix & evaluate your model's performance based on accuracy, precision, recall

Exercises

You will need to use the "caret" library. Create a .qmd-file and solve the tasks there. Store it in the JupyterHub folder "Session 10".

Out-of-sample prediction

Use your results from the logistic regression model from tutorial 9. Test how your model performs out-of-sample. Perform the following steps.

- 1. Split your data into a training and test sample (70% training, 30% testing).
- 2. Train your model on the training data.
- 3. Make predictions on the test set using the trained model.
- 4. Calculate the out-of-sample RMSE, i.e. the test set RMSE.

```
# clean environment
rm(list=ls())

# load packages/libraries

# install.packages("remotes") # hashtagged because already installed
library(remotes)
remotes::install_gitlab("BAQ6370/sozoekds", host="gitlab.rrz.uni-hamburg.de")
library(sozoekds)

library(tidyverse)
library(dplyr)
library(caret) # for splitting, CV
```

```
airbnb_data <- airbnbsmall # store data as "airbnb_data"</pre>
##################
# 1. split data into training and test set
# Define the x-variables of your model
xvar <- c("n_number_of_reviews", "price", "d_gym") # creates a column w/ those 3 vari
x <- airbnb_data[, xvar] # defines x as the variables in xvar</pre>
# Define the output variable of your model
airbnb_data$high_rating = ifelse(airbnb_data$n_review_scores_rating>94, 1, 0) # creat
y <- airbnb_data$high_rating # defines y as "high_rating" in the airbnb dataset
# Split the data into training and testing sets (70% training, 30% testing)
set.seed(123) # for reproducibility; to get the same random split each time you run
# creates the training data
trainIndex <- createDataPartition(y, p = 0.7, # percentage of data going to training
                                   list = FALSE,
                                   times = 1) # only 1 split
x_train <- x[trainIndex, ]</pre>
x test <- x[-trainIndex, ]</pre>
y_train <- y[trainIndex]</pre>
y_test <- y[-trainIndex]</pre>
# 2. Train your model on the training data
model <- glm(y_train ~ ., family=binomial(link="logit"), data = data.frame(cbind(y_tr</pre>
summary(model)
# 3. Make predictions on the test set using the trained model
out_predictions <- predict(model, newdata = data.frame(x_test), type="response") # ty</pre>
# 4. Calculate the out-of-sample RMSE, i.e. the test set RMSE.
residuals <- (y_test-out_predictions) # calculate the error (predicted values-observe
sqrt(mean(residuals^2)) # prints 0.4933515
```

Cross-validation, confusion matrix & evaluation metrics

Now apply the cross-validation procedure using the training data. Use the "trainControl" command and do a 5-fold and 10-fold cross-validation. Compare the results.

```
# specify model to be estimated using training data & apply cv
modelcross <- train(y train ~ ., data=data.frame(cbind(y train, x train)),</pre>
                method="glm",
                family=binomial,
                trControl=cross) # applies our cross-validation process as specified
modelcross # prints RMSE 0.4941984
### 10-fold CV ###
set.seed(123) # for reproducability
# specify cross-validation & number of folds
cross <- trainControl(method="cv", # cross-validation</pre>
                          number=10, # k=10
                      verboseIter = TRUE)
# specify model to be estimated using training data & apply cv
modelcross <- train(y_train ~ ., data=data.frame(cbind(y_train, x_train)),</pre>
                method="glm",
                family=binomial,
                trControl=cross) # applies our cross-validation process as specified
modelcross # prints RMSE 0.4946834
```

To evaluate your model create a confusion matrix with the "confusionMatrix" command and find accuracy, precision and recall scores.

For the confusion matrix you will need two classes "negative" and "positive" of the predicted values as well as two classes "negative" and "positive" of the observed values from the test set. Predicted probabilities <0.5 should fall in the predicted class "negative" (0), whereas predicted probabilities >0.5 should fall in the predicted class "positive" (1).

```
# create the classes of the predicted values and store in "p_class"
p_class <- factor(ifelse(out_predictions > 0.5, 1, 0)) # creates a factor being 1 whe

# levels in predicted values and observed values need to be the same, but are differe unique(levels(p_class)) # prints "Negative_Class" "Positive_Class" unique(levels(y_test)) # prints NULL

# therefore transform y_test variables into factor w/ same labels as p_class y_test_f <- factor(y_test, levels = c(1, 0))
tab <- table(predict = p_class, actual = y_test_f) # create table with predicted and confusionMatrix(p_class, y_test_f, mode = "prec_recall") # prints confusion matrix</pre>
```