Own_project

José Antonio Cardoso

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Introduction

To complete the Professional Data Science training, the student was asked to develop a project of his own (authored). Some rules were established, for example, that the purpose of the project object (algorithm) was freely chosen, as well as the data set to be used in this development. Therefore, this report presents the aforementioned algorithm, as well as describing its objective and functionality.

Overview

The project object (algorithm) uses classification techniques and for this, it uses Random Forest, a machine learning algorithm that uses decision trees to make predictions. It also uses a GUIDE data set made available by Microsoft, on the Kaggle platform, which is a global community of data scientists. (/kaggle/input/microsoft-security-incident-prediction/).

Executive Summary

The project objective (algorithm) is to classify cybersecurity incidents by analyzing the responses to calls made by the SOC (Security Operations Center) team, as described in the aforementioned data set.

Methods

Several R language packages are required to enable the development of the algorithm. The dataset is originally available on the Kaggle platform, but a copy of it was downloaded and made available together with the files for this project.

```
# Convert the Id field to character from the training dataset
GUIDE train$Id <-as.character(GUIDE train$Id)</pre>
# Convert the Id field to character from the test data set
GUIDE test$Id <-as.character(GUIDE test$Id)</pre>
# As you can see in this code, the data set is already divided.
# However, to demonstrate the knowledge about the logic and necessity of this
division
# I present here the code related to this process, where for educational
purposes I consider the
# complete GUDE file, which I fictitiously called complet GUIDE.csv
# Just for information...
# Note: The division between Training and Testing is 70% and 30% respectively
# Dividing the data into training and validation
# Defining the seed for generating random numbers
#set.seed(0)
#train index <- createDataPartition(complet GUIDE$IncidentGrade, p = 0.7,</pre>
list = FALSE)
#GUIDE train <- complet GUIDE[train index,]
#GUIDE_test <- complet_GUIDE[-train_index,]
```

Analysis

Originally the GUIDE dataset (Training and Testing) contains over 13 million pieces of evidence across 33 entity types, covering 1.6 million alerts and 1 million annotated incidents, distributed across 45 columns with information such as: DeviceId (Unique identifier for the device), IpAddress (Involved IP address), Url (Involved URL) AccountUpn (Email account identifier) etc. However, in order to meet the objective of classifying cybersecurity incidents by analyzing the responses to the calls made by the SOC (Security Operations Center) team, as described in the "Executive Summary", we decided to use the identification columns ("Id" and "IncidentId") as a matter of good practice, and we used the columns ActionGrouped (SOC alert remediation action (high level)), ActionGranular (SOC alert remediation action (fine grain)) and LastVerdict (Final verdict of the threat analysis) as predictor variables and IncidentGrade (SOC grade assigned to the incident) is the response variable.

```
c("Id", "IncidentId", "ActionGrouped", "ActionGranular", "LastVerdict", "IncidentG
rade")
# Apply the selection on the Training dataset
df_data_train <- df_data_train[, cols_df]</pre>
# Filters only the rows that contain SOC responses, thus eliminating any
column that contains "Not Available" values
df data train <- df data train %>%
   filter(ActionGrouped != "" & ActionGranular != "" & LastVerdict != "" &
IncidentGrade != "")
# Presents some quantitative information about the dataset.
cat("Number of records in the Training dataset","\n")
## Number of records in the Training dataset
cat("Show the first few rows of the dataset","\n")
## Show the first few rows of the dataset
head(df data train,5)
               Id IncidentId ActionGrouped ActionGranular LastVerdict
##
## 1 936302872206
                       56426 IsolateDevice isolateresponse Suspicious
## 2 627065227429
                      138497 IsolateDevice isolateresponse Suspicious
## 3 807453856769
                        8065 IsolateDevice isolateresponse Suspicious
                      327814 IsolateDevice quarantinefile Suspicious
## 4 970662611054
## 5 652835031931
                        5371 IsolateDevice isolateresponse Suspicious
##
      IncidentGrade
## 1
     TruePositive
## 2
      TruePositive
## 3
      TruePositive
## 4
      TruePositive
## 5 BenignPositive
cat("\n", "Show unique values from columns")
##
## Show unique values from columns
# Apply the function to each column and store the results in a list
newcols df <-
c("ActionGrouped", "ActionGranular", "LastVerdict", "IncidentGrade")
list result <- map(newcols df, ~df data train %>%
                  pull(.x) %>%
                  unique())
names(list_result) <- newcols_df</pre>
list result
## $ActionGrouped
## [1] "IsolateDevice" "ContainAccount"
```

```
##
## $ActionGranular
## [1] "isolateresponse"
## [2] "quarantinefile"
## [3] "change user password."
## [4] "disable account."
## [5] "update stsrefreshtokenvalidfrom timestamp."
## [6] "forcepasswordresetremediation"
## [7] "account password changed"
## [8] "disableuser"
## [9] "account disabled"
## [10] "reset user password."
## [11] "set force change user password."
## [12] "msecidentitiessuspenduser"
## [13] "msecidentitiesconfirmusercompromised"
## $LastVerdict
## [1] "Suspicious" "NoThreatsFound" "Malicious"
##
## $IncidentGrade
## [1] "TruePositive"
                      "BenignPositive" "FalsePositive"
# In this line above, the code prints the list containing the unique values
of
# each column, as explained above,
cat("\n","Shows the structure of the dataset","\n")
##
## Shows the structure of the dataset
str(df_data_train)
## 'data.frame': 1110 obs. of 6 variables:
                   : chr "936302872206" "627065227429" "807453856769"
## $ Id
"970662611054" ...
## $ IncidentId : int 56426 138497 8065 327814 5371 40983 119255 19426
766 5371 ...
## $ ActionGrouped : chr "IsolateDevice" "IsolateDevice" "IsolateDevice"
"IsolateDevice" ...
## $ ActionGranular: chr "isolateresponse" "isolateresponse"
"isolateresponse" "quarantinefile" ...
## $ LastVerdict : chr "Suspicious" "Suspicious" "Suspicious"
"Suspicious" ...
## $ IncidentGrade : chr "TruePositive" "TruePositive" "TruePositive"
"TruePositive" ...
cat("\n","Summarizes the dataset")
##
## Summarizes the dataset
```

```
summary(df data train)
##
         Id
                         IncidentId
                                        ActionGrouped
                                                            ActionGranular
##
   Length:1110
                       Min. : 14
                                        Length:1110
                                                            Length:1110
   Class :character
                       1st Qu.: 5371
##
                                        Class :character
                                                            Class :character
   Mode :character
                       Median : 31790
                                        Mode :character
                                                            Mode :character
##
                       Mean
                             : 72686
##
                       3rd Qu.: 96466
##
                              :566689
                       Max.
##
   LastVerdict
                       IncidentGrade
##
    Length:1110
                       Length:1110
   Class :character
                       Class :character
##
##
   Mode :character
                       Mode :character
##
##
##
# Transform the string column into a factor
df data train$IncidentGrade <- as.factor(df data train$IncidentGrade)</pre>
df data train$ActionGrouped <- as.factor(df data train$ActionGrouped)</pre>
df data train$ActionGranular <- as.factor(df data train$ActionGranular)</pre>
df_data_train$LastVerdict <- as.factor(df_data_train$LastVerdict)</pre>
```

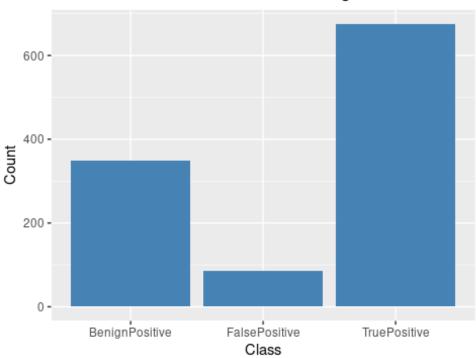
Execution

The model is executed using the machine learning algorithm that uses decision trees to make predictions, in this case Random Forest. It is extracted from the train function, which trains the model using Random Forest in an unencapsulated way. The trained_model object generated by the train function has the various methods that we use to demonstrate the execution of the model, including graphs of class distribution and importance of variables. The test is performed using the trained_model object, generating the appropriate predictions.

```
LastVerdict,
                       data = df data train,
                       method = "rf",
                       tuneGrid = hyper_grid,
                       trControl = ctrl)
# Print the trained model
print(trained_model)
## Random Forest
##
## 1110 samples
##
     3 predictor
      3 classes: 'BenignPositive', 'FalsePositive', 'TruePositive'
##
##
## No pre-processing
## Resampling: Cross-Validated (10 fold)
## Summary of sample sizes: 999, 998, 998, 1000, 999, 999, ...
## Resampling results across tuning parameters:
##
##
    mtry Accuracy
                     Kappa
          0.6874503 0.3256632
##
    3
##
    4
          0.6856485 0.3217928
##
    5
          0.6883756 0.3287985
##
## Accuracy was used to select the optimal model using the largest value.
## The final value used for the model was mtry = 5.
# Presents an overview of the different hyperparameter combinations tested
and their respective performance metrics.
trained_model$results
    mtry Accuracy
                       Kappa AccuracySD
                                            KappaSD
        3 0.6874503 0.3256632 0.02626770 0.05512794
## 1
## 2
       4 0.6856485 0.3217928 0.02640943 0.05542052
        5 0.6883756 0.3287985 0.02770996 0.05931271
## 3
# Show cross-validation results.
trained_model$resample
##
                    Kappa Resample
      Accuracy
## 1 0.6936937 0.3368477
                            Fold01
## 2 0.6339286 0.2283650
                            Fold02
## 3 0.6936937 0.3177874
                            Fold05
## 4 0.6909091 0.3352293
                            Fold04
## 5 0.6964286 0.3601075
                            Fold03
## 6 0.7027027 0.3444882
                            Fold06
## 7 0.7027027 0.3697522
                            Fold09
## 8 0.6756757 0.2806481
                            Fold08
     0.6576577 0.2722567
## 9
                            Fold07
## 10 0.7363636 0.4425026 Fold10
```

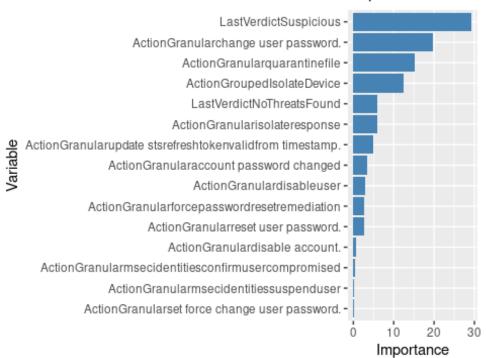
```
# Shows the final selected model after hyperparameter sweeping and cross-
validation.
trained_model$finalModel
##
## Call:
    randomForest(x = x, y = y, mtry = param$mtry)
                  Type of random forest: classification
##
                        Number of trees: 500
##
## No. of variables tried at each split: 5
##
           OOB estimate of error rate: 31.62%
##
## Confusion matrix:
                  BenignPositive FalsePositive TruePositive class.error
## BenignPositive
                                                         183
                             165
                                              0
                                                               0.5258621
## FalsePositive
                              32
                                              0
                                                          54
                                                               1.0000000
## TruePositive
                                                         594
                              82
                                                               0.1213018
# Shows the relative importance of each predictor variable in the final model
trained model$finalModel$importance
##
                                                             MeanDecreaseGini
## ActionGroupedIsolateDevice
                                                                   12.5354011
## ActionGranularaccount password changed
                                                                    3.5399873
## ActionGranularchange user password.
                                                                   19.6301176
## ActionGranulardisable account.
                                                                    0.7235116
## ActionGranulardisableuser
                                                                    2.9308146
## ActionGranularforcepasswordresetremediation
                                                                    2.7904143
## ActionGranularisolateresponse
                                                                    5.8971866
## ActionGranularmsecidentitiesconfirmusercompromised
                                                                    0.4269763
## ActionGranularmsecidentitiessuspenduser
                                                                    0.1364855
## ActionGranularquarantinefile
                                                                   15.2801257
## ActionGranularreset user password.
                                                                    2.5494853
## ActionGranularset force change user password.
                                                                    0.1045615
## ActionGranularupdate stsrefreshtokenvalidfrom timestamp.
                                                                    4.8546657
## LastVerdictNoThreatsFound
                                                                    5.9898430
## LastVerdictSuspicious
                                                                   29.1997856
# This graph shows the distribution of classes in the data set, helping to
understand if there is an imbalance.
# Plots the distribution of classes
ggplot(df_data_train, aes(x = IncidentGrade)) +
  geom bar(fill = "steelblue") +
  labs(title = "Distribution of Classes in the Training Set", x = "Class", y
= "Count")
```

Distribution of Classes in the Training Set



```
# This graph shows the relative importance of each predictor variable.
# Plots the importance of variables
importance <- varImp(trained_model, scale = FALSE)
ggplot(importance, aes(x = reorder(Overall, Overall), y = Overall)) +
    geom_bar(stat = "identity", fill = "steelblue") +
    coord_flip() +
    labs(title = "Importance of Variables", x = "Variable", y = "Importance")
### Coordinate system already present. Adding new coordinate system, which
will
### replace the existing one.</pre>
```

Importance of Va



```
# Make predictions on the TRAINING dataset to evaluate the model's
performance
train_predictions <- predict(trained_model, newdata = df_data_train)</pre>
# Converts categorical data into factors for classification
actual_class <- df_data_train$IncidentGrade</pre>
predicted_class <- train_predictions</pre>
# Calculates accuracy, as an assessment of model performance
accuracy_train <- accuracy(actual_class, predicted_class)</pre>
cat("Accuracy:", accuracy_train, "\n")
## Accuracy: 0.6963964
# Calculate the F1-score, as an evaluation of the model's performance
f1_train <- f1(actual_class, predicted_class)</pre>
cat("F1-score:", f1_train, "\n")
## F1-score: 0.8
# TEST AND EVALUATE THE TRAINED MODEL
# Make a copy of the test dataset
df_data_test <- GUIDE_test</pre>
```

```
# Apply the selection to the Test dataset
df_data_test <- df_data_test[, cols_df]</pre>
# Filters only the rows that contain SOC responses, thus eliminating any
column that contains "Not Available" values
df data test <- df data test %>%
   filter(ActionGrouped != "" & ActionGranular != "" & LastVerdict != "" &
IncidentGrade != "")
# Transform the string column into a factor
df data test$IncidentGrade <- as.factor(df data test$IncidentGrade)</pre>
df data test$ActionGrouped <- as.factor(df data test$ActionGrouped)</pre>
df_data_test$ActionGranular <- as.factor(df_data_test$ActionGranular)</pre>
df data_test$LastVerdict <- as.factor(df_data_test$LastVerdict)</pre>
# Make predictions on the TEST dataset to evaluate the model's performance
test_predictions <- predict(trained_model, newdata = df_data_test)</pre>
# Converts categorical data into factors for classification
actual class <- df data test$IncidentGrade</pre>
predicted_class <- test_predictions</pre>
# Calculates accuracy, as an assessment of model performance
accuracy test <- accuracy(actual class, predicted class)</pre>
cat("Accuracy:", accuracy_test, "\n")
## Accuracy: 0.7058824
# Calculate the F1-score, as an evaluation of the model's performance
f1 test <- f1(actual class, predicted class)
cat("F1-score:", f1_test, "\n")
## F1-score: 0.8
```

Results

Although the data set is considerably large, which is very good for achieving greater reliability in the model's execution, I tried to eliminate as many rows and columns of incomplete data as possible, thus generating a smaller amount of data, but with all rows and columns containing reliable information. I tried to use accuracy to evaluate the model's performance, as well as the F1 Score to help balance the importance of false positives and false negatives.

```
## Accuracy: 0.6963964

# Calculate the F1-score, as an evaluation of the model's performance
cat("F1-score:", f1_train, "\n")

## F1-score: 0.8

# Calculates accuracy, as an assessment of model performance
cat("Accuracy:", accuracy_test, "\n")

## Accuracy: 0.7058824

# Calculate the F1-score, as an evaluation of the model's performance
cat("F1-score:", f1_test, "\n")

## F1-score: 0.8
```

Conclusion

Reaching an accuracy of 0.7058824 and an F1-Score of 0.8 can be considered a good result in some aspects, but I understand that there is a lot of room for improvement. The exclusion of incomplete data rows and columns, as mentioned above, contributed to obtaining these performance numbers, although we were left with a more limited amount of data. However, in order to continue the work, I intend to reevaluate the data set, looking for more subsidies for the model to make its classifications.