Milestone 4: Model Analysis

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```
library(readr)
library(ggplot2)
library(gridExtra)
library(dplyr)
##
## Attaching package: 'dplyr'
## The following object is masked from 'package:gridExtra':
##
##
       combine
## The following objects are masked from 'package:stats':
##
       filter, lag
##
## The following objects are masked from 'package:base':
       intersect, setdiff, setequal, union
##
library(ggcorrplot)
library(fastDummies)
library(corrplot)
## corrplot 0.94 loaded
library(randomForest)
## randomForest 4.7-1.2
## Type rfNews() to see new features/changes/bug fixes.
##
## Attaching package: 'randomForest'
## The following object is masked from 'package:dplyr':
##
##
       combine
## The following object is masked from 'package:gridExtra':
##
##
## The following object is masked from 'package:ggplot2':
##
##
       margin
```

```
library(caret)

## Loading required package: lattice
library(caret)

setwd('C:/Users/nosip/Documents/third Year/BIN381/milestones')
```

Model Assessment: Random Forest Model

The following Milestone will be dedicated to the tuning of the already trained random forest model with the aim of improving the accuracy and precision of the model.

###Read the data Read in the data that has been prepared for the random forest (rf) model:

```
[4] "Annual_Salary"
                              "Months_Annual"
                                                    "FRS.Contribution"
##
   [7] "Net_months"
                              "Gross_Salary"
                                                    "Gross_Months"
## [10] "Qualify"
                              "Education_Bach."
                                                    "Education_HS_grad"
## [13] "Education_Masters"
                                                    "Occupation_Exec."
                              "Occupation_Cleric."
## [16] "Occupation Prof."
                              "Occupation Sales"
                                                    "age"
```

View the structure to ensure that the data types are as expected (only numeric).

```
str(data_df)
```

```
151133 obs. of 18 variables:
##
  'data.frame':
   $ marital_status
##
                       : int
                             1 2 2 1 2 2 1 2 2 2 ...
##
   $ household_size
                       : int
                             2 2 2 2 2 2 2 2 2 2 ...
                             4 4 4 4 4 4 4 4 4 ...
## $ yrs_of_residence : int
## $ Annual Salary
                             0.62 0.25 0.394 0.735 0.386 ...
                       : num
                              2 3 4 1 6 3 18 2 4 8 ...
##
   $ Months Annual
                       : int
## $ FRS.Contribution : num 0.617 0.223 0.501 0.433 0.949 ...
## $ Net_months
                       : num -0.516 -0.211 -0.482 -0.958 2.335 ...
                       : num 0.0255 0.9321 0.9684 0.4696 0.8508 ...
## $ Gross_Salary
                       : num -0.539 -0.192 -0.47 -0.956 2.31 ...
## $ Gross Months
## $ Qualify
                       : int 0000111000...
## $ Education_Bach.
                       : int 0000000000...
## $ Education_HS_grad : int
                             0 0 0 0 0 0 0 0 0 0 ...
## $ Education_Masters : int
                             1 1 1 1 1 1 1 1 1 1 . . .
## $ Occupation_Cleric.: int
                             0 0 0 0 0 0 0 0 0 0 ...
## $ Occupation_Exec. : int
                             0 0 0 0 0 0 0 0 0 0 ...
   $ Occupation Prof.
                             1 1 1 1 1 1 1 1 1 1 ...
                       : int
##
   $ Occupation_Sales
                      : int
                              0 0 0 0 0 0 0 0 0 0 ...
                       : int
                             48 60 82 47 75 74 78 46 75 76 ...
```

The data types for the columns are as expected. The Gross Months and Net Months have been previously normalised and scaled; hence some values are negative; this is to be expected.

The histograms below show the columns before they were scaled:

```
data_b4_scaling <- read.csv("data_for_ml.csv", header = TRUE)</pre>
```

```
columns_to_distr <- c("Annual_Salary", "FRS.Contribution",</pre>
"Net_months", "Gross_Salary", "Gross_Months")
# folr loop to loop through the above columns:
plot_list <- list()</pre>
for (column in columns_to_distr) {
p <- ggplot(data_b4_scaling, aes_string(x = column)) +</pre>
geom histogram(binwidth = 50, fill = "turquoise", color = "black", alpha = 0.7) +
labs(title = paste("Distribution of", column), x = column, y = "Frequency") +
theme minimal()
plot_list[[column]] <- p</pre>
}
## Warning: `aes_string()` was deprecated in ggplot2 3.0.0.
## i Please use tidy evaluation idioms with `aes()`.
## i See also `vignette("ggplot2-in-packages")` for more information.
## This warning is displayed once every 8 hours.
## Call `lifecycle::last_lifecycle_warnings()` to see where this warning was
## generated.
grid.arrange(grobs = plot_list, ncol = 2)
                                                          Distribution of FRS.Contribution
          Distribution of Annual_Salary
                                                  Erednency
6000
6000
2000
0
Frequency
    10000
     5000
        0
                                                        0
            0
                   250
                           500
                                   750
                                           1000
                                                             0
                                                                    250
                                                                             500
                                                                                     750
                                                                                             1000
                     Annual_Salary
                                                                     FRS.Contribution
          Distribution of Net_months
                                                          Distribution of Gross_Salary
                                                  Evednency 8000 8000 8000 9000 9000 9000 0
 Frequency
   75000
   50000
   25000
        0
             0
                      100
                                200
                                          300
                                                             0
                                                                    250
                                                                                     750
                                                                             500
                                                                                             1000
                      Net_months
                                                                       Gross_Salary
           Distribution of Gross_Months
    100000
 Frequency
     75000
     50000
     25000
         0
              0
                       100
                                          300
                                 200
                      Gross_Months
```

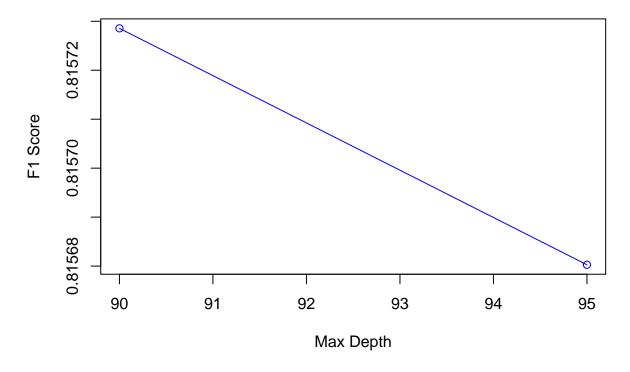
However in milestone 3, the data was scaled appropriately.

Split the data

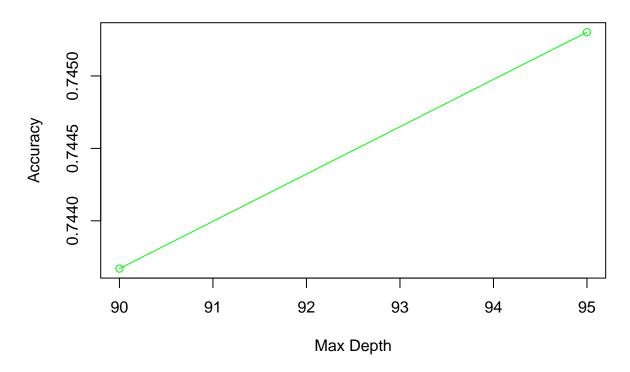
The data will be split as following: - Training - Testing - Validation

```
#split the data
set.seed(123)
total_rows <- nrow(data_df)</pre>
#split data 70-30
train_indices <- sample(1:total_rows, 0.7 * total_rows)</pre>
train_data <- data_df[train_indices, ]</pre>
remaining_indices <- setdiff(1:total_rows, train_indices)</pre>
#testing and validation will each make up 15%
validation_indices <- sample(remaining_indices, 0.5 * length(remaining_indices))</pre>
test_indices <- setdiff(remaining_indices, validation_indices)</pre>
validation_data <- data_df[validation_indices, ]</pre>
test_data <- data_df[test_indices, ]</pre>
cat("Training data size:", nrow(train_data), "\n")
## Training data size: 105793
cat("Validation data size:", nrow(validation_data), "\n")
## Validation data size: 22670
cat("Testing data size:", nrow(test_data), "\n")
## Testing data size: 22670
The rf Model (Max Depth Tuning)
The first parameter that will be tuned is the max-depth parameter of the tree; this will restrict how deep the
trees go and this parameter will help with reducing over fitting.
formula <- Qualify ~ marital_status + household_size + yrs_of_residence +
Annual_Salary + Months_Annual + FRS.Contribution + Net_months + Gross_Salary + Gross_Months +
Education_Bach. + Education_HS_grad + Education_Masters +
Occupation_Cleric. + Occupation_Exec. + Occupation_Prof. +
Occupation_Sales + age
#metrices
depth_values <- c(90, 95)
accuracy_scores <- c()</pre>
f1_scores <- c()</pre>
convert Qualify to factors; because in its numeric form the model assumes a regression nature.
train_data$Qualify <- as.factor(train_data$Qualify)</pre>
validation_data$Qualify <- as.factor(validation_data$Qualify)</pre>
test_data$Qualify <- as.factor(test_data$Qualify)</pre>
for (depth in depth_values) {
  # Train the random forest model with the current depth
  model <- randomForest(formula, data = train_data, maxnodes = depth)</pre>
  # Make predictions on the validation dataset
  predictions <- predict(model, newdata = validation_data)</pre>
  # Convert predictions to factors to match the validation data's Qualify column
  predictions <- as.factor(predictions)</pre>
  # Calculate the confusion matrix
```

F1 Score at Different Depths



Accuracy at Different Depths



The tree depth was cycled from 0 to 95. As the depth increases; it increases the accuracy score and F1 score. However, this is a slow increase; hence the next attempt will be to tune the features from the previously generated Gini Feature importance from Milestone 3.

Feature Tuning (Gini Importance)

```
names(data_df)
    [1] "marital_status"
                              "household_size"
                                                     "yrs_of_residence"
##
    [4]
        "Annual_Salary"
                              "Months_Annual"
                                                    "FRS.Contribution"
    [7] "Net_months"
                              "Gross_Salary"
                                                    "Gross Months"
   [10] "Qualify"
##
                              "Education_Bach."
                                                     "Education_HS_grad"
       "Education_Masters"
                              "Occupation_Cleric."
                                                    "Occupation_Exec."
   [13]
                                                    "age"
   [16] "Occupation_Prof."
                              "Occupation_Sales"
```

from the above Gini importance the following columns appear to have little impact on the model; hence they will be dropped: - marital_status - yrs_of_residence - Occupation_Sales - Education_Bach. - Education_Masters - Occupation_Prof. - household_size - Education_HS_grad - Occupation_Exec. - Occupation_Exec.

Drop columns with low Gini importance

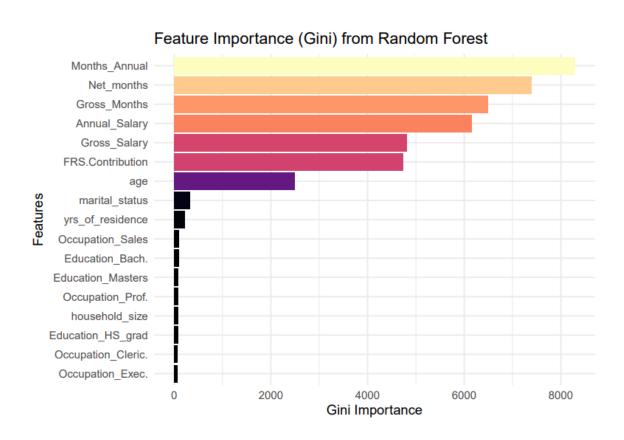


Figure 1: Ft importance

Split Data

```
#split the data
set.seed(123)
total_rows <- nrow(data_df)</pre>
#split data 70-30
train_indices <- sample(1:total_rows, 0.7 * total_rows)</pre>
train_data <- data_df[train_indices, ]</pre>
remaining_indices <- setdiff(1:total_rows, train_indices)</pre>
#testing and validation will each make up 15%
validation_indices <- sample(remaining_indices, 0.5 * length(remaining_indices))</pre>
test_indices <- setdiff(remaining_indices, validation_indices)</pre>
validation_data <- data_df[validation_indices, ]</pre>
test_data <- data_df[test_indices, ]</pre>
define formula without the columns:
# Updated formula
formula <- Qualify ~ Annual_Salary + Months_Annual + FRS.Contribution + Net_months + Gross_Salary + Gro
Occupation_Cleric. + age
```

```
formula <- Qualify ~ Annual_Salary + Months_Annual + FRS.Contribution + Net_months + Gross_Occupation_Cleric. + age

#metrices
depth_values <- c(40, 45)
accuracy_scores <- c()
f1_scores <- c()
```

convert Qualify to factors; because in its numeric form the model assumes a regression nature.

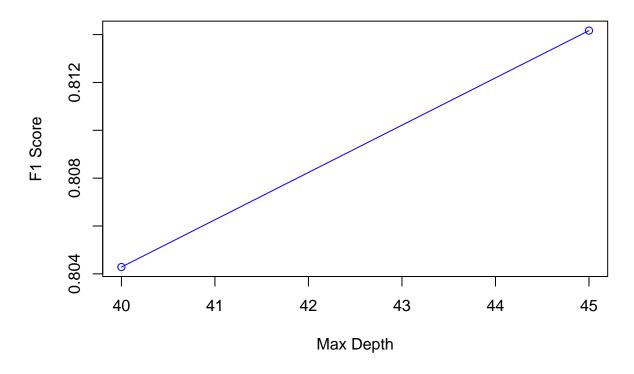
```
train_data$Qualify <- as.factor(train_data$Qualify)
validation_data$Qualify <- as.factor(validation_data$Qualify)
test_data$Qualify <- as.factor(test_data$Qualify)</pre>
```

Train the model again

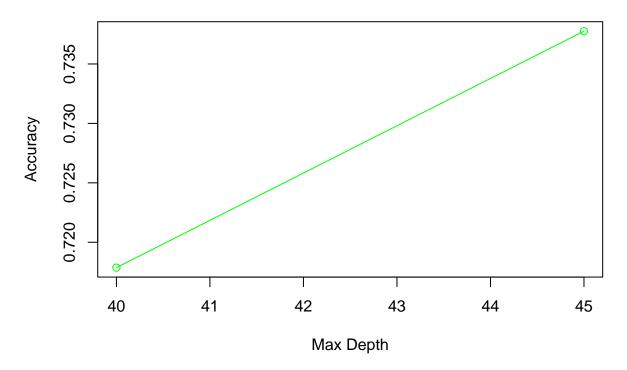
```
for (depth in depth values) {
  # Train the random forest model with the current depth
 model <- randomForest(formula, data = train_data, maxnodes = depth)</pre>
  # Make predictions on the validation dataset
  predictions <- predict(model, newdata = validation_data)</pre>
  # Convert predictions to factors to match the validation data's Qualify column
  predictions <- as.factor(predictions)</pre>
  # Calculate the confusion matrix
  confusion_matrix <- confusionMatrix(predictions, validation_data$Qualify)</pre>
  # Extract Accuracy from confusion matrix
  accuracy_scores <- c(accuracy_scores, confusion_matrix$overall['Accuracy'])</pre>
  # Extract F1 score from confusion matrix (using 'F1' method from caret)
 f1_scores <- c(f1_scores, confusion_matrix$byClass['F1'])</pre>
}
# Plot F1 Score vs Max Depth
plot(depth_values, f1_scores, type = "o", col = "blue",
```

```
xlab = "Max Depth", ylab = "F1 Score",
main = "F1 Score at Different Depths")
```

F1 Score at Different Depths



Accuracy at Different Depths



The F1 score and accuracy increased; up until a max depth of 40 (accuracy = 71%; f1-score = 80%); after this accuracy consistently decreased; while the f1-score increased. The Aim is to strike a balance between these two metrices.

Number of Trees Parameter

The next parameter to tune is the Number of trees parameter; along with the depth.

```
depth_values <- c(80, 85)
trees_values <- c(150, 200)

accuracy_scores <- list()
f1_scores <- list()

# Loop through each depth
for (depth in depth_values) {

# Store metrics for each depth
accuracy_depth <- c()
f1_depth <- c()

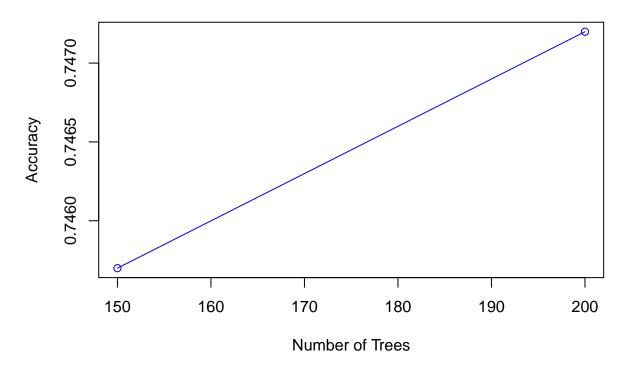
# Loop through each number of trees
for (num_trees in trees_values) {

# Train the Random Forest model with current depth and number of trees
model <- randomForest(formula, data = train_data, maxnodes = depth, ntree = num_trees)

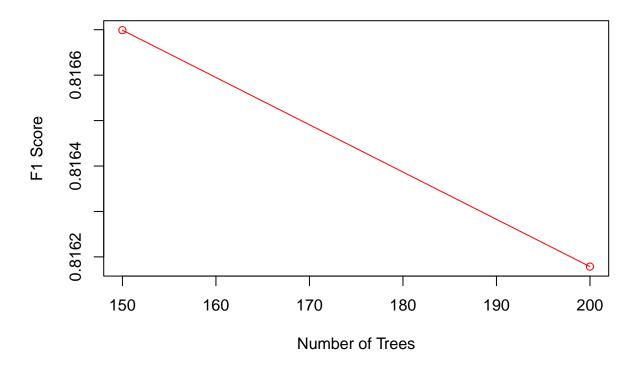
# Make predictions on the validation dataset</pre>
```

```
predictions <- predict(model, newdata = validation_data)</pre>
  # Calculate confusion matrix
  confusion_matrix <- confusionMatrix(predictions, validation_data$Qualify)</pre>
  # Extract accuracy and F1 score
 accuracy_depth <- c(accuracy_depth, confusion_matrix$overall['Accuracy'])</pre>
 f1 depth <- c(f1 depth, confusion matrix$byClass['F1'])
}
# Store the accuracy and F1 scores for this depth
accuracy_scores[[as.character(depth)]] <- accuracy_depth</pre>
f1 scores[[as.character(depth)]] <- f1 depth</pre>
# Plot accuracy vs. number of trees for this depth
plot(trees_values, accuracy_depth, type = "o", col = "blue",
     xlab = "Number of Trees", ylab = "Accuracy",
     main = paste("Accuracy vs. Number of Trees (Depth =", depth, ")"))
# Plot F1 score vs. number of trees for this depth
plot(trees_values, f1_depth, type = "o", col = "red",
     xlab = "Number of Trees", ylab = "F1 Score",
     main = paste("F1 Score vs. Number of Trees (Depth =", depth, ")"))
```

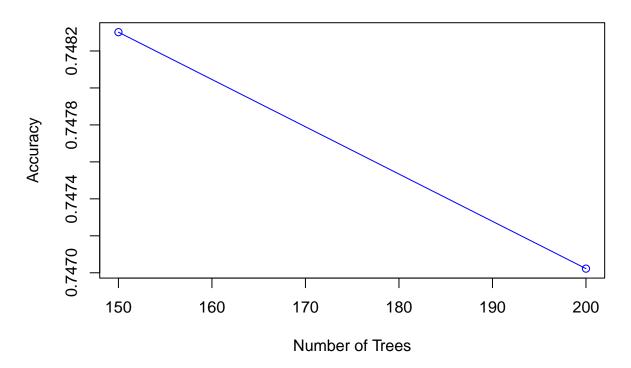
Accuracy vs. Number of Trees (Depth = 80)

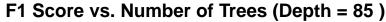


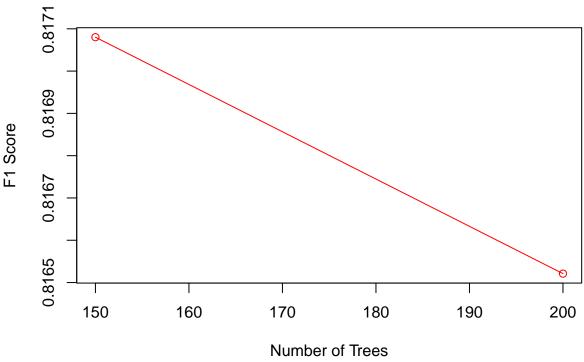
F1 Score vs. Number of Trees (Depth = 80)



Accuracy vs. Number of Trees (Depth = 85)







The highest accuracy score is 74% at a depth of 85; and a number of trees of 150. This also correlated with the highest f1 score of 82%. However; these do not beat the original untuned model that will be given below.

Original untuned Model

```
model_retrained <- randomForest(Qualify ~ ., data = train_data)

# Validate the model again using the validation set
validation_predictions_retrained <- predict(model_retrained, newdata = validation_data)

validation_confusion_matrix_retrained <- table(Actual = validation_data$Qualify, Predicted = validation

performance metrics:

accuracy_retrained <- sum(diag(validation_confusion_matrix_retrained)) / sum(validation_confusion_matrix_
sensitivity_retrained <- validation_confusion_matrix_retrained[2, 2] / sum(validation_confusion_matrix_
specificity_retrained <- validation_confusion_matrix_retrained[1, 1] / sum(validation_confusion_matrix_
precision_retrained <- validation_confusion_matrix_retrained[2, 2] / sum(validation_confusion_matrix_retrained)

fl_score_retrained <- 2 * (precision_retrained * recall_retrained) / (precision_retrained + recall_retr

cat("Accuracy:", round(accuracy_retrained * 100, 2), "%\n")

## Accuracy: 96.9 %

cat("Sensitivity (Recall):", round(sensitivity_retrained * 100, 2), "%\n")

## Sensitivity (Recall): 95.88 %
```

```
cat("Specificity:", round(specificity_retrained * 100, 2), "%\n")

## Specificity: 97.59 %

cat("Precision:", round(precision_retrained * 100, 2), "%\n")

## Precision: 96.37 %

cat("F1 Score:", round(f1_score_retrained * 100, 2), "%\n")

## F1 Score: 96.12 %
```

Metrics Interpretation

Accuracy (88.04%): - The model correctly predicted 87% of the total records in the validation set. The classes are slightly imbalanced but not significantly as to highly affect the accuracy score.

- Recall (Sensitivity) (81.5%):
 - The model's ability to correctly classify positive (1) records. The model correctly identified 81% of True positives.
 - Which means 18.5% of true positives we predicted to be negative.
 - The model correctly identified 81.5% of customers who qualify for the service.
- Specificity (92.25%):
 - The models ability to classify true negatives.
 - the model successfully identified 95% of applicants who do not qualify for the service.
- Precision (87.76%):
 - This is the measure of the accuracy of positive predictions.
- F1 Score (84.51%): This is the true measure of the performance of the model. it is the harmonic mean of precision and sensitivity. An 84% F1-Score indicates a solid model performance in predicting customer who qualify for the service and those who do not.

```
print(names(data_df))
```

The original model performs better than the tuned model. hence we will be continuing with the original model.

```
## [1] "Annual_Salary" "Months_Annual" "FRS.Contribution"
## [4] "Net_months" "Gross_Salary" "Gross_Months"
## [7] "Qualify" "Occupation Cleric." "age"
```

Save updated model

```
saveRDS(model_retrained, "retrained_rf_model2.rds")
```

Conclusion

The tuning of the Random Forest model through adjusting both depth and number of trees yielded the highest accuracy of 74% at a depth of 85 with 150 trees. This also correlated with the highest F1 score of 82%. These improvements indicate that careful tuning can lead to better performance of the model.

However; the original un-tuned model; with the newly dropped column that have a low feature importance score; still outperformed the tuned model. Hence, the original model will be carried on to the next milestones.