LogisticRegression_Model.R

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```
# Load required libraries for Risk Assessment
library(tidyverse)
## Warning: package 'tidyverse' was built under R version 4.3.3
## Warning: package 'ggplot2' was built under R version 4.3.3
## Warning: package 'tidyr' was built under R version 4.3.3
## Warning: package 'dplyr' was built under R version 4.3.3
## Warning: package 'stringr' was built under R version 4.3.3
## Warning: package 'forcats' was built under R version 4.3.3
## Warning: package 'lubridate' was built under R version 4.3.3
## -- Attaching core tidyverse packages ------ tidyverse 2.0.0 --
## v dplyr 1.1.4
                      v readr
                                   2.1.5
## v forcats 1.0.0 v stringr 1.5.1
## v ggplot2 3.5.1
                    v tibble
                                   3.2.1
## v lubridate 1.9.3
                       v tidyr
                                   1.3.1
## v purrr
              1.0.2
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                    masks stats::lag()
## i Use the conflicted package (<a href="http://conflicted.r-lib.org/">http://conflicted.r-lib.org/</a>) to force all conflicts to become error
library(caret)
## Warning: package 'caret' was built under R version 4.3.3
## Loading required package: lattice
## Attaching package: 'caret'
```

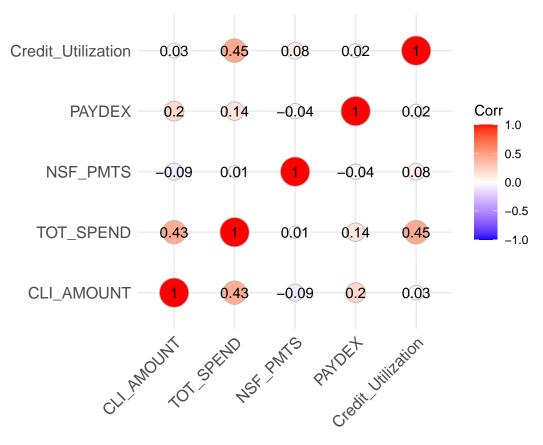
The following object is masked from 'package:purrr':

##

lift

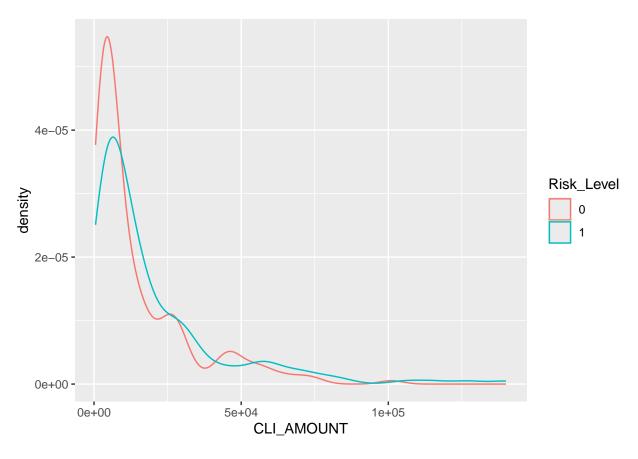
```
library(pROC)
## Warning: package 'pROC' was built under R version 4.3.3
## Type 'citation("pROC")' for a citation.
##
## Attaching package: 'pROC'
## The following objects are masked from 'package:stats':
##
       cov, smooth, var
library(car)
## Warning: package 'car' was built under R version 4.3.3
## Loading required package: carData
## Warning: package 'carData' was built under R version 4.3.3
##
## Attaching package: 'car'
## The following object is masked from 'package:dplyr':
##
##
       recode
##
## The following object is masked from 'package:purrr':
##
##
       some
library(ggcorrplot)
## Warning: package 'ggcorrplot' was built under R version 4.3.3
library(readxl)
## Warning: package 'readxl' was built under R version 4.3.3
# Load dataset
data <- read_excel("ACCT_Monitoring_FinalData.xlsx")</pre>
# Data Preprocessing
# 1. Handle missing values
data <- data %>% mutate(across(where(is.numeric), ~ifelse(is.na(.), mean(., na.rm = TRUE), .)))
# 2. Calculate derived feature for Credit Utilization
data <- data %>% mutate(Credit_Utilization = TOT_SPEND / CREDIT_LIMIT)
```

```
# 3. Create a target variable for risk assessment
# High risk is defined as Credit Utilization > 0.8
data <- data %>% mutate(Risk_Level = ifelse(Credit_Utilization > 0.8, 1, 0))
# 4. Select relevant features
selected_features <- c("CLI_AMOUNT", "TOT_SPEND", "NSF_PMTS", "PAYDEX",</pre>
                        "Credit_Utilization", "Risk_Level")
data <- data %>% select(all_of(selected_features))
# Ensure target variable is a factor
data$Risk_Level <- as.factor(data$Risk_Level)</pre>
\# Train-Test Split
set.seed(123)
train_index <- createDataPartition(data$Risk_Level, p = 0.7, list = FALSE)
train_data <- data[train_index, ]</pre>
test_data <- data[-train_index, ]</pre>
# Correlation:
cor_matrix <- cor(train_data %>% select(where(is.numeric)))
print(cor_matrix)
##
                       CLI AMOUNT
                                    TOT SPEND
                                                   NSF PMTS
                                                                 PAYDEX
## CLI AMOUNT
                       1.00000000 0.434303486 -0.092004516 0.20045405
## TOT_SPEND
                       0.43430349 1.000000000 0.005135212 0.13964078
## NSF_PMTS
                      -0.09200452 0.005135212 1.000000000 -0.03743348
## PAYDEX
                       0.20045405 0.139640783 -0.037433485 1.00000000
## Credit Utilization 0.02645997 0.446951872 0.078267738 0.01983681
##
                      Credit_Utilization
## CLI_AMOUNT
                              0.02645997
## TOT_SPEND
                              0.44695187
## NSF_PMTS
                              0.07826774
                              0.01983681
## PAYDEX
## Credit_Utilization
                              1.00000000
# Visualize correlations
ggcorrplot(cor_matrix, method = "circle", lab = TRUE)
```



```
# Fit a logistic regression model
model_glm <- glm(Risk_Level ~ ., data = train_data, family = "binomial", control = glm.control(maxit =</pre>
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
# Calculate VIF for logistic regression model
vif_results <- vif(model_glm)</pre>
print(vif_results)
##
           CLI_AMOUNT
                                TOT_SPEND
                                                     NSF_PMTS
                                                                           PAYDEX
##
             4.723723
                                 4.785640
                                                     1.012916
                                                                         1.051277
## Credit_Utilization
             1.078458
# Predict on the test data
pred_glm <- predict(model_glm, newdata = test_data, type = "response")</pre>
# Evaluate with ROC and AUC
roc_glm <- roc(test_data$Risk_Level, pred_glm)</pre>
## Setting levels: control = 0, case = 1
## Setting direction: controls < cases
```

```
print(paste("AUC for Logistic Regression (GLM): ", auc(roc_glm)))
## [1] "AUC for Logistic Regression (GLM): 1"
# Model Summary
summary(model_glm)
##
## Call:
## glm(formula = Risk_Level ~ ., family = "binomial", data = train_data,
       control = glm.control(maxit = 1000))
##
## Coefficients:
##
                        Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                     -4.422e+02 5.436e+05 -0.001
                                                      0.999
## CLI_AMOUNT
                                             0.000
                                                      1.000
                     -2.027e-04 5.850e+00
                      1.572e-03 8.168e+00
## TOT_SPEND
                                             0.000
                                                      1.000
                      -5.385e+00 1.107e+06
## NSF_PMTS
                                             0.000
                                                      1.000
## PAYDEX
                       2.580e-02 4.050e+03
                                             0.000
                                                      1.000
## Credit_Utilization 5.181e+02 5.189e+05
                                             0.001
                                                      0.999
## (Dispersion parameter for binomial family taken to be 1)
##
##
       Null deviance: 4.8556e+02 on 350 degrees of freedom
## Residual deviance: 1.1976e-09 on 345 degrees of freedom
## AIC: 12
##
## Number of Fisher Scoring iterations: 34
ggplot(train_data, aes(x = CLI_AMOUNT, color = Risk_Level)) + geom_density()
```

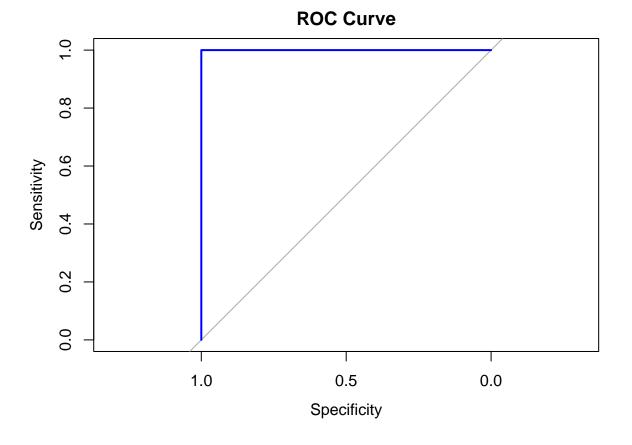


```
# Predictions
pred_probs <- predict(model_glm, test_data, type = "response")
pred_classes <- ifelse(pred_probs > 0.5, 1, 0)

# Performance Metrics
conf_matrix <- confusionMatrix(factor(pred_classes), test_data$Risk_Level)
print(conf_matrix)</pre>
```

```
## Confusion Matrix and Statistics
##
##
             Reference
## Prediction 0 1
            0 79 0
##
##
            1 0 70
##
##
                  Accuracy : 1
                    95% CI : (0.9755, 1)
##
##
       No Information Rate : 0.5302
       P-Value [Acc > NIR] : < 2.2e-16
##
##
                     Kappa: 1
##
##
##
    Mcnemar's Test P-Value : NA
##
##
               Sensitivity: 1.0000
               Specificity: 1.0000
##
```

```
##
            Pos Pred Value : 1.0000
##
            Neg Pred Value : 1.0000
                Prevalence: 0.5302
##
##
            Detection Rate: 0.5302
      Detection Prevalence : 0.5302
##
##
         Balanced Accuracy: 1.0000
##
          'Positive' Class : 0
##
##
# AUC-ROC
roc_curve <- roc(as.numeric(test_data$Risk_Level), pred_probs)</pre>
## Setting levels: control = 1, case = 2
## Setting direction: controls < cases</pre>
auc <- auc(roc_curve)</pre>
plot(roc_curve, col = "blue", main = "ROC Curve")
```



```
print(paste("AUC:", auc))
```

[1] "AUC: 1"