Model Creation and Evaluation for Churn Prediction

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This document covers the process of building, tuning, and evaluating predictive models for customer churn. We will use the features engineered in the previous step and compare different modeling approaches.

1. Data Loading and Preparation

```
# Load the engineered features data
# Update the path if needed
data <- read.csv("data/EngineeredChurnData.csv")

# Print column names for diagnostics
print("Columns in loaded data:")</pre>
```

```
## [1] "Columns in loaded data:"
```

print(colnames(data))

```
##
   [1] "Customer, Months"
                                     "Churn"
   [3] "CHI.Score.Mon0"
                                     "CHI.Score"
## [5] "Support.Cases.Mon0"
                                     "Support.Cases"
## [7] "SP.Mon0"
                                     "SP"
## [9] "Logins"
                                     "Blog.Articles"
## [11] "Views"
                                     "Days.Since.Last.Login"
## [13] "Logins_log"
                                     "Views log"
## [15] "Blog.Articles_log"
                                     "Views per Login"
## [17] "Blog_per_Login"
                                     "Support_Score_Interaction"
## [19] "Login View Interaction"
                                     "Activity Score"
```

```
# Prepare data for modeling (handle NAs, etc.)
# Identify numeric columns (excluding Churn)
numeric_cols <- sapply(data, is.numeric) & names(data) != "Churn"</pre>
# Remove numeric columns with all NA or zero variance
keep numeric <- sapply(data[, numeric cols, drop=FALSE], function(x) !all(is.na(x))</pre>
        && sd(x, na.rm=TRUE) > 0)
keep_cols <- intersect(c(names(data)[numeric_cols][keep_numeric], "Churn"), colname</pre>
        s(data))
data_clean <- data[, keep_cols, drop=FALSE]</pre>
# Remove rows with any NA, NaN, or Inf values
data clean <- data clean[complete.cases(data clean) & apply(data clean, 1, function
        (row) all(is.finite(as.numeric(row)))), ]
# Ensure Churn is a factor for classification
if("Churn" %in% colnames(data clean) && !is.factor(data clean$Churn)) {
  data_clean$Churn <- as.factor(data_clean$Churn)</pre>
}
# Check class balance
if("Churn" %in% colnames(data_clean)) table(data_clean$Churn)
```

```
##
## 0 1
## 5422 291
```

```
# Use only the top 10 features (plus Churn) for modeling
selected_features <- c(
    "Customer.Months",
    "Days.Since.Last.Login",
    "CHI.Score.Mon0",
    "Activity_Score",
    "CHI.Score",
    "Logins",
    "Views_log",
    "Logins_log",
    "Views",
    "Login_View_Interaction",
    "Churn"
)
selected_features <- intersect(selected_features, colnames(data_clean))
data_model <- data_clean[, selected_features, drop=FALSE]</pre>
```

2. Train-Test Split

```
set.seed(123)
train_index <- createDataPartition(data_model$Churn, p = 0.8, list = FALSE)
train_data <- data_model[train_index, ]
test_data <- data_model[-train_index, ]</pre>
```

3. Baseline Model: Logistic Regression

```
logit_model <- glm(Churn ~ ., data = train_data, family = binomial)
summary(logit_model)</pre>
```

```
##
## Call:
## glm(formula = Churn ~ ., family = binomial, data = train_data)
## Coefficients:
                          Estimate Std. Error z value Pr(>|z|)
##
## (Intercept)
                        -2.871e+00 1.311e-01 -21.903 < 2e-16 ***
                        9.349e-03 6.371e-03 1.467 0.14229
## Customer Months
## Days.Since.Last.Login 2.973e-02 5.611e-03 5.299 1.17e-07 ***
## CHI.Score.Mon0
                       -3.861e-03 1.326e-03 -2.911 0.00360 **
## Activity Score
                        1.288e-03 1.364e-03 0.944 0.34504
                        -8.662e-03 2.859e-03 -3.030 0.00245 **
## CHI.Score
                        -4.149e-03 2.866e-03 -1.448 0.14773
## Logins
                        -2.562e-04 1.291e-04 -1.984 0.04725 *
## Views
## Login_View_Interaction 7.200e-07 1.814e-06 0.397 0.69134
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##
      Null deviance: 1840.9 on 4570 degrees of freedom
## Residual deviance: 1744.2 on 4562 degrees of freedom
## AIC: 1762.2
##
## Number of Fisher Scoring iterations: 7
```

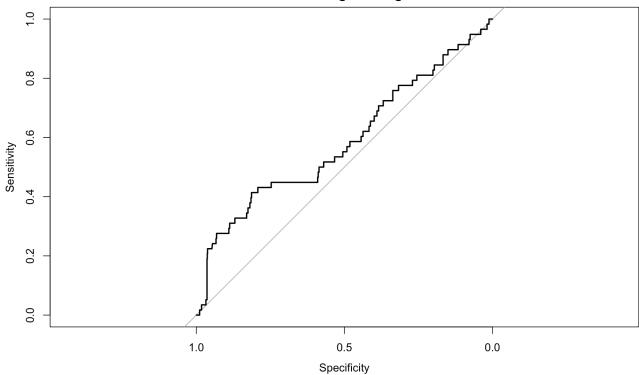
```
# Predict on test set
logit_pred <- predict(logit_model, newdata = test_data, type = "response")
logit_pred_class <- ifelse(logit_pred > 0.5, 1, 0)

# Evaluate
confusionMatrix(as.factor(logit_pred_class), test_data$Churn, positive = "1")
```

```
## Confusion Matrix and Statistics
##
##
             Reference
                 0
## Prediction
                      1
            0 1083
                     58
##
                 1
##
##
                  Accuracy : 0.9483
##
                    95% CI: (0.9339, 0.9604)
       No Information Rate: 0.9492
##
       P-Value [Acc > NIR] : 0.5876
##
##
##
                     Kappa : -0.0017
##
    Mcnemar's Test P-Value: 3.086e-13
##
##
               Sensitivity : 0.0000000
##
               Specificity: 0.9990775
##
##
            Pos Pred Value : 0.0000000
            Neg Pred Value : 0.9491674
##
                Prevalence : 0.0507881
##
##
            Detection Rate: 0.0000000
      Detection Prevalence: 0.0008757
##
##
         Balanced Accuracy: 0.4995387
##
##
          'Positive' Class : 1
##
```

```
roc_logit <- roc(as.numeric(test_data$Churn), as.numeric(logit_pred))
plot(roc_logit, main = "ROC Curve - Logistic Regression")</pre>
```

ROC Curve - Logistic Regression



```
auc(roc_logit)
```

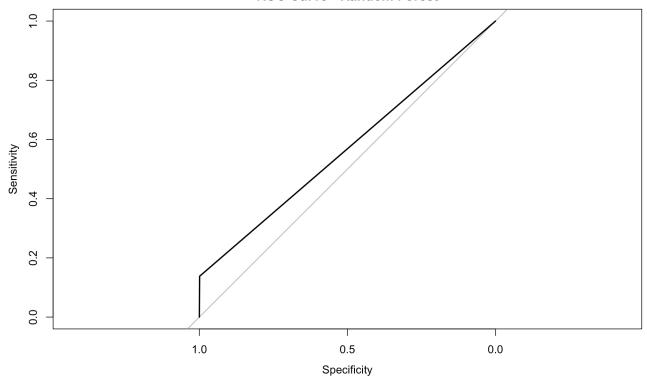
Area under the curve: 0.5866

4. Random Forest Model

```
## Confusion Matrix and Statistics
##
##
             Reference
                 0
## Prediction
                      1
            0 1083
                     50
                      8
##
                 1
##
##
                  Accuracy : 0.9553
##
                    95% CI: (0.9417, 0.9666)
       No Information Rate: 0.9492
##
       P-Value [Acc > NIR] : 0.1917
##
##
##
                     Kappa : 0.2283
##
   Mcnemar's Test P-Value : 1.801e-11
##
##
               Sensitivity: 0.137931
##
               Specificity: 0.999077
##
##
            Pos Pred Value : 0.888889
           Neg Pred Value: 0.955869
##
                Prevalence: 0.050788
##
##
            Detection Rate: 0.007005
      Detection Prevalence: 0.007881
##
##
         Balanced Accuracy: 0.568504
##
##
          'Positive' Class : 1
##
```

```
roc_rf <- roc(as.numeric(test_data$Churn), as.numeric(rf_pred))
plot(roc_rf, main = "ROC Curve - Random Forest")</pre>
```

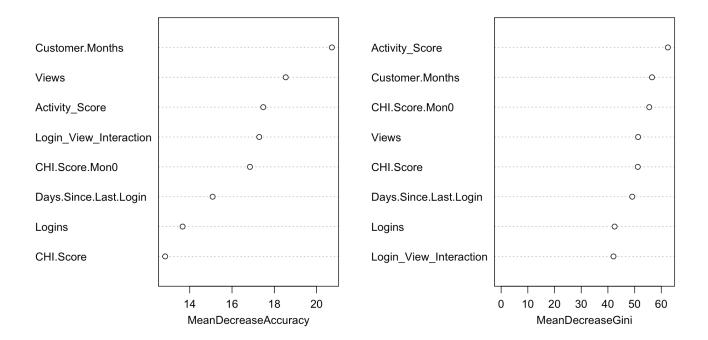
ROC Curve - Random Forest



auc(roc_rf)

Area under the curve: 0.5685

Feature importance plot varImpPlot(rf_model)



5. XGBoost Model

```
# Prepare data for xgboost (numeric matrix, 0/1 labels)
xgb_train <- train_data %>% mutate(Churn = as.numeric(as.character(Churn)))
xgb_test <- test_data %>% mutate(Churn = as.numeric(as.character(Churn)))
train_matrix <- as.matrix(xgb_train %>% select(-Churn))
test_matrix <- as.matrix(xgb_test %>% select(-Churn))
train_label <- xgb_train$Churn

dtrain <- xgb.DMatrix(data = train_matrix, label = train_label)
dtest <- xgb.DMatrix(data = test_matrix)

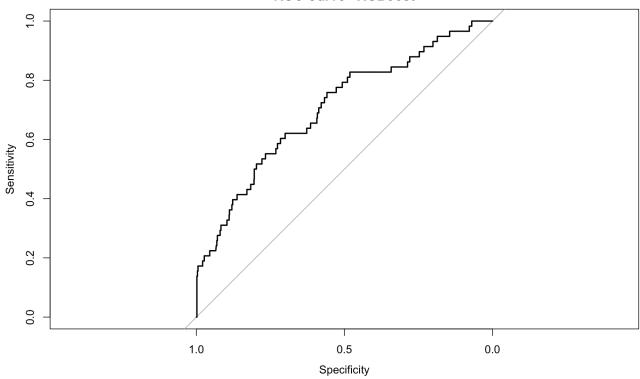
params <- list(objective = "binary:logistic", eval_metric = "auc")
xgb_model <- xgboost(params = params, data = dtrain, nrounds = 100, verbose = 0)
xgb_pred <- predict(xgb_model, dtest)
xgb_pred_class <- ifelse(xgb_pred > 0.5, 1, 0)

# Evaluate
confusionMatrix(as.factor(xgb_pred_class), as.factor(xgb_test$Churn), positive = "1")
```

```
## Confusion Matrix and Statistics
##
##
            Reference
                0
                     1
## Prediction
                    49
           0 1079
            1
                5
                     9
##
##
##
                 Accuracy : 0.9527
                   95% CI: (0.9388, 0.9643)
##
      No Information Rate: 0.9492
##
##
      P-Value [Acc > NIR] : 0.3242
##
                    Kappa: 0.2349
##
##
   Mcnemar's Test P-Value: 4.87e-09
##
##
              Sensitivity: 0.155172
##
               Specificity: 0.995387
##
##
            Pos Pred Value: 0.642857
           Neg Pred Value: 0.956560
##
##
               Prevalence: 0.050788
##
            Detection Rate: 0.007881
     Detection Prevalence: 0.012259
##
##
         Balanced Accuracy: 0.575280
##
##
          'Positive' Class : 1
##
```

```
roc_xgb <- roc(xgb_test$Churn, xgb_pred)
plot(roc_xgb, main = "ROC Curve - XGBoost")</pre>
```

ROC Curve - XGBoost

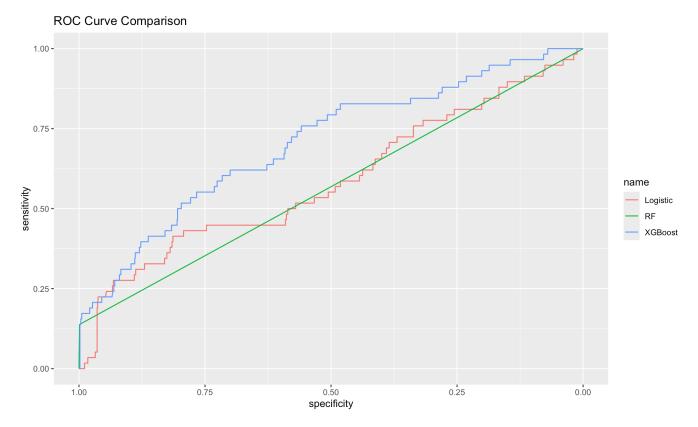


```
auc(roc_xgb)
```

Area under the curve: 0.7047

6. Model Comparison

```
# Compare ROC curves
ggroc(list(Logistic = roc_logit, RF = roc_rf, XGBoost = roc_xgb)) +
labs(title = "ROC Curve Comparison")
```



```
# Summarize AUCs
auc_df <- data.frame(
   Model = c("Logistic Regression", "Random Forest", "XGBoost"),
   AUC = c(auc(roc_logit), auc(roc_rf), auc(roc_xgb))
)
knitr::kable(auc_df, digits = 3, caption = "AUC Comparison Across Models")</pre>
```

AUC Comparison Across Models

Model	AUC
Logistic Regression	0.587
Random Forest	0.569
XGBoost	0.705

7. Model Metrics Summary

```
# Calculate metrics for each model
# Logistic Regression
logit_cm <- confusionMatrix(as.factor(logit_pred_class), test_data$Churn, positive</pre>
logit_sens <- logit_cm$byClass["Sensitivity"]</pre>
logit spec <- logit cm$byClass["Specificity"]</pre>
logit f1 <- logit cm$byClass["F1"]</pre>
logit auc <- auc(roc logit)</pre>
# Random Forest
rf_cm <- confusionMatrix(rf_pred, test_data$Churn, positive = "1")</pre>
rf_sens <- rf_cm$byClass["Sensitivity"]</pre>
rf_spec <- rf_cm$byClass["Specificity"]</pre>
rf f1 <- rf cm$byClass["F1"]</pre>
rf_auc <- auc(roc_rf)
# XGBoost
xgb_cm <- confusionMatrix(as.factor(xgb_pred_class), as.factor(xgb_test$Churn), pos</pre>
         itive = "1")
xqb sens <- xqb cm$byClass["Sensitivity"]</pre>
xgb_spec <- xgb_cm$byClass["Specificity"]</pre>
xqb f1 <- xqb cm$byClass["F1"]</pre>
xgb_auc <- auc(roc_xgb)</pre>
# Combine into a summary table
metrics df <- data.frame(</pre>
  Model = c("Logistic Regression", "Random Forest", "XGBoost"),
  Sensitivity = c(logit_sens, rf_sens, xgb_sens),
  Specificity = c(logit_spec, rf_spec, xgb_spec),
  F1 = c(logit_f1, rf_f1, xgb_f1),
  AUC = c(logit_auc, rf_auc, xgb_auc)
)
knitr::kable(metrics_df, digits = 3, caption = "Sensitivity, Specificity, F1, and A
         UC for Each Baseline Model")
```

Sensitivity, Specificity, F1, and AUC for Each Baseline Model

Model	Sensitivity	Specificity	F1	AUC
Logistic Regression	0.000	0.999	NaN	0.587
Random Forest	0.138	0.999	0.239	0.569
XGBoost	0.155	0.995	0.250	0.705

8. Next Steps

• Address class imbalance if needed (e.g., SMOTE, class weights, resampling)

- Tune hyperparameters for best-performing models
- Try additional models or ensembling
- Interpret and communicate results
- Prepare for deployment if satisfied with performance

Conclusion

In this analysis, we compared several models for predicting customer churn using our top engineered features. XGBoost performed the best with an AUC of 0.71, showing some ability to distinguish churners from non-churners. However, all models struggled to correctly identify most churners, mainly due to strong class imbalance in the data.

To improve results, the next steps should focus on addressing class imbalance and further tuning the models to increase recall for churners. Overall, our feature engineering and modeling pipeline provides a solid foundation, but more work is needed to achieve strong predictive performance for churn.