ML approach

Decision trees frequently perform well on imbalanced data. In modern machine learning, tree ensembles, such as Random Forest, almost always outperform singular decision trees. Random Forests is selected because its performance compared with other machine learning techniques is the best in terms of classification metrics and low upper bound error probability. The analysis is explained in details in the following sections. The appendix contains complete code to:

- Create train and test sets.
- Create a Random Forest classifier (including setting class weights).
 To combat severe class imbalance of datasets, Random Forest with class weighting for imbalanced classification will be used.
- Tune Hyperparameter by Grid Search
 The exhaustive search for the optimal number of trees and the minimum samples of leaf is used. The minimum samples of leaf of all models for all the datasets are all set 2. The number of trees for Dataset a is 125, while those for other datasets are 250.
- Evaluate the model using two metrics (accuracy and AUC)
- Evaluate the mode's upper bound probability of misclassification

Methodology for validation

To begin with the ML-based classification, 80 percent of each dataset was used for training, with 5-fold cross-validation, and 20 percent of the datasets has been used for testing. Both linear and nonlinear classifiers have been tested based on their classification metrics and upper bound probability to select the best ML model. Random Forest (RF) will be compared with other nonlinear models, including Penalized Support Vector Machine (SVM), XGBoost (XGB), and Neural Network (NN), and a linear model, Logistic regression (LR), accompanied by SMOTE resampling. The accuracy and AUC are used to measure the classification performance of each model.

Methodology for upper bound estimation for the misclassification error

The probability of the error or misclassification can be calculated with (1) [1]

$$P(error) = \int_{-\infty}^{+\infty} P(error|x) P(x) dx$$

If we estimate the probability density function of each class, by dividing the error probability into two regions as R_1 and R_2 , the probability of error can be written as:

$$\int_{\mathbb{R}_{1}} P(x|Class\ 0)\ P(Class\ 0)\ dx + \int_{\mathbb{R}_{2}} P(x|Class\ 1)\ P(Class\ 1)\ dx$$

Fukunaga, K. (1990) showed that the upper bound error can be calculated using probability density function (PDF)-based distances, which usually rely on mean and variance distance. For

safety concern, it is critical to consider the worst-case scenario. In this case, Chernoff Distance (CD), as a Probability Density Function (PDF)-based distance measure, is selected to test an upper bound for the probability of the next misclassification error [2].

$$P(error) = P(Class\ 0)^{\lambda}\ P(Class\ 1)^{1-\lambda}$$

$$\int_{-\infty}^{+\infty} P(Class\ 0)^{\lambda}\ P(Class\ 1)^{1-\lambda}\ dx$$

Dataset a: Table 1 compares the upper bound error probability estimation based on Chernoff Distance. We observe that XGB model reports the lowest upper bound, followed by RF classifier.

Table 1: Comparison of Chernoff Distance for Dataset a

	LR	SVM	RF	XGB	NN
CD	0. 116673	0.032613	0.018716	0.008076	0.020670

Table 2 represents the accuracy and AUC of each ML model for Dataset a. RF model has the best accuracy and AUC overall.

Table 2: Comparison of classification metrics for Dataset a

	LR	SVM	RF	XGB	NN
Accuracy	0.485	0.895	0.945	0.925	0.87
AUC	0.483	0.878	0.931	0.905	0.914

Dataset b: Similar to the Dataset a, the proposed Chernoff Distance can be applied to the Dataset b. Seeing from Table 3, SVM and RF models have better estimation for upper bound probability for the larger Dataset b.

Table 3: Comparison of Chernoff Distance for Dataset b

	LR	SVM	RF	XGB	NN
CD	0. 002134	0.000859	0.001198	0.001317	0.001770

Table 4 represents the accuracy and AUC of each ML model for Dataset b. RF model has the best accuracy and AUC overall.

Table 4: Comparison of classification metrics for Dataset b

	LR	SVM	RF	XGB	NN
Accuracy	0.859	0.905	0.989	0.982	0.859
AUC	0.860	0.906	0.989	0.982	0.952

Dataset c: The Dataset c has similar statistical characteristics with the Dataset b. Table 5 provides the upper bound error probability of different classifiers. As indicated in the table, the best estimation is related to SVM model. The following RF and XGB models also have relatively low upper bound probability.

Table 5: Comparison of Chernoff Distance for Dataset c

	LR	SVM	RF	XGB	NN
CD	0. 115958	0.000660	0.001186	0.001110	0.010075

Table 6 represents the accuracy and AUC of each ML models for Dataset c. RF model has the best accuracy and AUC overall.

Table 6: Comparison of classification metrics for Dataset c

	LR	SVM	RF	XGB	NN
Accuracy	0.619	0.557	0.994	0.974	0.818
AUC	0.566	0.728	0.983	0.971	0.930

Overall, it is noticeable that RF have the best classification performance and relatively low Chernoff Distance, so it is selected as a reliable safety prediction model.

Scalability to Higher Dimension

The Chernoff Distance can be calculated in a higher dimension as long as mean and variance distances of every dimension is provided. In order to load high-dimension data into ML models, the solution to this flexibility should be considered. The two feasible approaches to adapt ML models to higher dimension are:

- Increase the size of input layer (limited to Neural Network): since it provides flexibility to
 adjust input layer without changing hidden layer and output layer, it could be a simple
 way to allow the model to receive higher dimension of data. However, the model could
 not be a perfect fit to the new dimension of data since the data can have more noises
 than before.
- Dimension reduction: reduce the dimension of data before loaded into the model. This
 is an efficient way to identify the most important latent variable from the original data.
 However, concerning safety, we aim to catch as much information as possible. Tiny
 information loss, particularly about danger detection, can lead to an accident.

Reference

- [1] S. Theodoridis and K. Koutroumbas, Pattern Recognition. Elsevier Inc., 2009.
- [2] K. Fukunaga, Introduction to statistical pattern recognition. Elsevier, 2013.

Appendix

The following Python code is accessible on my Github

https://github.com/cory1219/Siemens-Challenge-/blob/main/Siemens%20final%20script.py

```
import os
import tempfile
import random
import pandas as pd
import numpy as np
import seaborn as sns
import matplotlib.pyplot as plt
from sklearn.model selection import train test split
from sklearn.model selection import cross val score
from sklearn.model selection import GridSearchCV
from imblearn.over_sampling import SMOTE
from sklearn.linear model import LogisticRegression
from sklearn.svm import SVC
from sklearn.ensemble import RandomForestClassifier
from xgboost import XGBClassifier
import tensorflow as tf
from tensorflow import keras
from tensorflow.python.keras import backend as K
from sklearn.metrics import confusion matrix
from sklearn.metrics import plot confusion matrix
from sklearn.metrics import accuracy_score, roc_auc_score
from bayes_opt import BayesianOptimization
#import datasets
random.seed(2021)
df1 = pd.read excel('trainingdata a.xls')
df2 = pd.read excel('trainingdata b.xls')
df3 = pd.read_excel('trainingdata_c.xls')
#create functions of spliting data for LOG, SVM, RF, XGBoost
def split(df):
  X_df = df[['x_i1', 'x_i2']]
  y df = df[['l i']]
  X_train, X_test, y_train, y_test = train_test_split(
    X df, y df, test size=0.2, random state=2021, stratify = y df)
  return X train, X test, y train, y test, y df
X_train_1, X_test_1, y_train_1, y_test_1, y_df_1 = split(df1) #dataset a
X_train_2, X_test_2, y_train_2, y_test_2, y_df_2 = split(df2) #dataset b
X_train_3, X_test_3, y_train_3, y_test_3, y_df_3 = split(df3) #dataset c
```

```
#Random forest
def rf(X train, y train, X test, y test):
  rf = RandomForestClassifier(class weight='balanced') #class weighting
  params rf = {'n estimators': [25, 100, 125, 250],
         'min samples leaf': [2, 10, 20],
         'random state': [2021]}
  grid rf = GridSearchCV(estimator=rf,
              param grid=params rf,
              scoring='roc auc',
              cv=5,
              verbose=1) #hyperparameter tuning
  grid rf.fit(X train, y train) #train
  print(grid rf.best score )
  best rf = grid rf.best estimator
  y_pred_rf = best_rf.predict(X_test) #predict
  plot confusion matrix(best rf, X test, y test)
  accuracy = accuracy score(y test, y pred rf) #accuracy
  auc = roc auc score(y test, y pred rf) #auc
  parameter = grid rf.best params
  print("Accuracy",accuracy score(y test, y pred rf))
  print("ROCAUC",roc auc score(y test, y pred rf))
  return y pred rf, accuracy, auc, parameter
y_pred_rf_1, accuracy_rf_1, auc_rf_1, parameter_rf_1 = rf(X_train_1, y_train_1, X_test_1,
y test 1) #dataset a
y pred rf 2, accuracy rf 2, auc rf 2, parameter rf 2 = rf(X train 2, y train 2, X test 2,
y test 2) #dataset b
y_pred_rf_3, accuracy_rf_3, auc_rf_3, parameter_rf_3 = rf(X_train_3, y_train_3, X_test_3,
y test 3) #dataset c
#PDF-based upper bounds
def chernoff distance(s, means, variances, univariate = False):
  """ Returns the Chernoff Distance, as defined in (3.150), p.98 of
    Fukunaga, 1990, Introduction to Statistical Pattern Recognition, 2nd Edition
  111111
  from math import log
  from math import sqrt
  from math import pow
  from numpy.linalg import det
  from numpy.linalg import inv
  from numpy import array
  if univariate:
```

```
part1 = 0.25 * log(0.25 * (variances[0] / variances[1] + variances[1] / variances[0]) + 0.5)
    part2 = 0.25 * pow(means[0] - means[1], 2) / (variances[0] + variances[1])
    return part1 + part2
  mean diff = array(means[0]) - array(means[1])
  var_avg = (array(variances[0]) + array(variances[1])) * 0.5
  In coeff = 0.5 * log(det(var avg) / sqrt(det(array(variances[0])) * det(array(variances[1]))))
  return 0.125 * np.matmul(np.matmul(mean diff.T, inv(var avg)), mean diff) + ln coeff
def upper bound(X train, y train, X test, y pred):
  return chernoff distance(
  0.5,
  [(X train[y train.l i == 1].x i1.mean(), X train[y train.l i == 1].x i2.mean()),
  (X_{test[y\_pred == 1].x_i1.mean(), X_{test[y\_pred == 1].x_i2.mean())],
  [[[X train[y train.l i == 1].x i1.var(), 0], [0, X train[y train.l i == 1].x i2.var()]],
  [[X \text{ test}[y \text{ pred} == 1].x \text{ i1.var}(), 0], [0, X \text{ test}[y \text{ pred} == 1].x \text{ i2.var}()]]])
#upper bound of RF
upper_bound(X_train_1, y_train_1, X_test_1, y_pred_rf_1) #dataset a
upper bound(X train 2, y train 2, X test 2, y pred rf 2) #dataset b
upper_bound(X_train_3, y_train_3, X_test_3, y_pred_rf_3) #dataset c
#comparison with other ML models
#Logistic regression with SMOTE
def log smote(X train, y train, X test, y test):
  sm = SMOTE(random state=2021, sampling strategy = "minority")
  X train, y_train = sm.fit_resample(X_train, y_train)
  print(y train.l i.sum())
  Ir sm = LogisticRegression(random state=2021).fit(X train, y train)
  y pred log = lr sm.predict(X test)
  #y pred prob = Ir sm.predict proba(X test)[:,1]
  plot_confusion_matrix(lr_sm, X_test, y_test)
  accuracy = accuracy score(y test, y pred log)
  auc = roc_auc_score(y_test, y_pred_log)
  print("Accuracy",accuracy score(y test, y pred log))
  print("ROCAUC",roc_auc_score(y_test, y_pred_log))
  return y pred log, accuracy, auc
y pred log 1, accuracy log 1, auc log 1 = log smote(X train 1, y train 1, X test 1,
y test 1)
y pred log 2, accuracy log 2, auc log 2 = log smote(X train 2, y train 2, X test 2,
y_test_2)
```

```
y_pred_log_3, accuracy_log_3, auc_log_3 = log_smote(X_train_3, y_train_3, X_test_3,
y test 3)
upper_bound(X_train_1, y_train_1, X_test_1, y_pred_log_1) #upper bound of log for dataset a
upper bound(X train 2, y train 2, X test 2, y pred log 2) #upper bound of log for dataset b
upper_bound(X_train_3, y_train_3, X_test_3, y_pred_log_3) #upper bound of log for dataset c
#Penalized SVM
def svm_pen(X_train, y_train, X_test, y_test):
  svm p = SVC(class weight = 'balanced', probability=True)
  svm p.fit(X train, y train)
  y pred svm = svm p.predict(X test)
  plot confusion matrix(svm p, X test, y test)
  accuracy = accuracy score(y test, y pred svm)
  auc = roc_auc_score(y_test, y_pred_svm)
  print("Accuracy",accuracy score(y test, y pred svm))
  print("ROCAUC",roc auc score(y test, y pred svm))
  return y_pred_svm, accuracy, auc
y pred svm 1, accuracy svm 1, auc svm 1 = svm pen(X train 1, y train 1, X test 1,
y test 1)
y_pred_svm_2, accuracy_svm_2, auc_svm_2 = svm_pen(X_train_2, y_train_2, X_test_2,
y_test 2)
y_pred_svm_3, accuracy_svm_3, auc_svm_3 = svm_pen(X_train_3, y_train_3, X_test_3,
y test 3)
upper bound(X train 1, y train 1, X test 1, y pred svm 1) #upper bound of svm for dataset
upper bound(X train 2, y train 2, X test 2, y pred svm 2) #upper bound of svm for dataset
upper_bound(X_train_3, y_train_3, X_test_3, y_pred_svm_3) #upper bound of svm for dataset
#XGBoost
def xgb(X train, y train, X test, y test, y df):
  est = (y df[y df['| i'] == 0].count() / y df[y df['| i'] == 1].count()).to numpy().item()
  xgb = XGBClassifier(scale pos weight=est)
  params xgb = {'n estimators': [3, 5, 10],
         'num_boost_round': [5, 10, 15],
         'max depth': [2, 5, 10],
         'eta': [0.001, 0.01, 0.1],
         'random state': [2021]}
  grid xgb = GridSearchCV(estimator=xgb,
             param grid=params xgb,
```

```
scoring='roc auc',
             cv=5,
             verbose=1)
  grid xgb.fit(X train, y train)
  print(grid xgb.best score )
  best xgb = grid xgb.best estimator
  y pred xgb = best xgb.predict(X test)
  plot confusion matrix(best xgb, X test, y test)
  accuracy = accuracy_score(y_test, y_pred_xgb)
  auc = roc auc score(y test, y pred xgb)
  print("Accuracy",accuracy score(y test, y pred xgb))
  print("ROCAUC",roc_auc_score(y_test, y_pred_xgb))
  return y pred xgb, accuracy, auc
y_pred_xgb_1, accuracy_xgb_1, auc_xgb_1 = xgb(X_train_1, y_train_1, X_test_1, y_test_1,
y df 1)
y pred xgb 2, accuracy xgb 2, auc xgb 2 = xgb(X train 2, y train 2, X test 2, y test 2,
y df 2)
y pred xgb 3, accuracy xgb 3, auc xgb 3 = xgb(X train 3, y train 3, X test 3, y test 3,
y df 3)
upper bound(X_train_1, y_train_1, X_test_1, y_pred_xgb_1) #upper bound of xgb for dataset a
upper bound(X train 2, y train 2, X test 2, y pred xgb 2) #upper bound of xgb for dataset b
upper_bound(X_train_3, y_train_3, X_test_3, y_pred_xgb_3) #upper bound of xgb for dataset c
#Neural networks weighted
tf.random.set seed(2021)
def separate(df): #create train, validation, test datasets
  neg, pos = np.bincount(df['l i'])
  total = neg + pos
  print('Examples:\n Total: {}\n Positive: {} ({:.2f}%)\n'.format(
    total, pos, 100*pos/total))
  cleaned df = df.copy()
  train df, test df = train test split(cleaned df, test size=0.2, random state=2021,
stratify=cleaned df['l i'])
  base df, val df = train test split(train df, test size=0.2, random state=2021,
stratify=train df['l i'])
  train labels = np.array(base df.pop('l i'))
  val labels = np.array(val df.pop('l i'))
  test labels = np.array(test df.pop('l i'))
  train features = np.array(base df[['x i1', 'x i2']])
  val features = np.array(val df[['x i1', 'x i2']])
  test features = np.array(test df[['x_i1', 'x_i2']])
```

```
initial bias = np.log([pos/neg])
  train_df_labels = train_df[['l_i']]
  weight for 0 = (1/neg)*(total)/2.0
  weight for 1 = (1/pos)*(total)/2.0
  class weight = {0: weight for 0, 1: weight for 1}
  class weight
  print('Weight for class 0: {:.2f}'.format(weight for 0))
  print('Weight for class 1: {:.2f}'.format(weight for 1))
  return train df, test df, train_labels, train_features, val_features, val_labels, test_labels,
test features, initial bias, class weight, train df labels
train df 1, test df 1, train labels 1, train features 1, val features 1, val labels 1,
test_labels_1, test_features_1, initial_bias_1, class_weight_1, train_df_labels_1 = separate(df1)
#dataset a
train df 2, test df 2, train labels 2, train features 2, val features 2, val labels 2,
test labels 2, test features 2, initial bias 2, class weight 2, train df labels 2 = separate(df2)
#dataset b
train df 3, test df 3, train labels 3, train features 3, val features 3, val labels 3,
test labels 3, test features 3, initial bias 3, class weight 3, train df labels 3 = separate(df3)
#dataset c
early stopping = tf.keras.callbacks.EarlyStopping(
  monitor='val auc',
  verbose=1,
  patience=10,
  mode='max',
  restore best weights=True)
def plot cm(labels, predictions, p=0.5): #confusion matrix plot
 cm = confusion matrix(labels, predictions > p)
 plt.figure(figsize=(5,5))
 sns.heatmap(cm, annot=True, fmt="d")
 plt.title('Confusion matrix @{:.2f}'.format(p))
 plt.ylabel('Actual label')
 plt.xlabel('Predicted label')
 print('Legitimate Transactions Detected (True Negatives): ', cm[0][0])
 print('Legitimate Transactions Incorrectly Detected (False Positives): ', cm[0][1])
 print('Fraudulent Transactions Missed (False Negatives): ', cm[1][0])
 print('Fraudulent Transactions Detected (True Positives): ', cm[1][1])
 print('Total Fraudulent Transactions: ', np.sum(cm[1]))
```

```
def nn(train df, test df, train labels, train features, val features, val labels, test labels,
test features, initial bias, class weight): #classifier
  def get model opt(units=16, dropout=0.5):
    output bias = tf.keras.initializers.Constant(initial bias)
    model = keras.Sequential([
    keras.layers.Dense(int(units),
               activation='relu',
               input shape=(train features.shape[-1],)),
    keras.layers.Dropout(dropout),
    keras.layers.Dense(1, activation='sigmoid', bias initializer=output bias)])
    return model
  def fit with(units, dropout, learning rate, epochs, batch size):
    model = get model opt(units, dropout)
    model.compile(
      optimizer=keras.optimizers.Adam(learning rate=learning rate),
      loss=keras.losses.BinaryCrossentropy(),
      metrics=[
        keras.metrics.BinaryAccuracy(name='accuracy'),
        keras.metrics.AUC(name='auc')
        1)
    initial weights = os.path.join(tempfile.mkdtemp(), 'initial weights')
    model.save weights(initial weights)
    model.load weights(initial weights)
    history = model.fit(x=train features,
               y=train labels,
               epochs=int(epochs),
               batch size=int(batch size),
               validation data=(val features,val labels),
               callbacks=[early stopping],
               class weight=class weight)
    # Evaluate the model with the eval dataset.
    accuracy = history.history['accuracy'][-1]
    K.clear session()
    tf.compat.v1.reset default graph()
    # Return the accuracy.
    return accuracy
  # Bounded region of parameter space
```

```
pbounds = {
  'units': (16,512),
  'dropout': (0.1,0.8),
  'batch size': (32, 128),
  'epochs': (25, 100),
  'learning rate': (1e-3, 1e-1)}
optimizer = BayesianOptimization(
f=fit with,
pbounds=pbounds,
verbose=2,
random state=2021
)
optimizer.maximize(init_points=10, n_iter=10,)
for i, res in enumerate(optimizer.res):
  print("Iteration {}: \n\t{}".format(i, res))
print(optimizer.max)
def get nnw model(units, dropout, learning rate, epochs, batch size):
  model = get_model_opt(units, dropout)
  model.compile(
    optimizer=keras.optimizers.Adam(learning rate=learning rate),
    loss=keras.losses.BinaryCrossentropy(),
    metrics=[
      keras.metrics.BinaryAccuracy(name='accuracy'),
      keras.metrics.AUC(name='auc')
      1)
  initial_weights = os.path.join(tempfile.mkdtemp(), 'initial_weights')
  model.save weights(initial weights)
  model.load weights(initial weights)
  model.fit(x=train features,
       y=train labels,
       epochs=int(epochs),
       batch size=int(batch size),
       validation data=(val features,val labels),
       callbacks=[early stopping],
       class_weight=class_weight)
  return model
```

```
model weighted = get nnw model(
    optimizer.max['params']['units'],
    optimizer.max['params']['dropout'],
    optimizer.max['params']['learning rate'],
    optimizer.max['params']['epochs'],
    optimizer.max['params']['batch size']
  train predictions baseline weighted = np.where(model weighted.predict(train features,
batch size=128)>=0.5, 1, 0)
  test predictions baseline weighted = np.where(model weighted.predict(test features,
batch size=128)>=0.5, 1, 0)
  baseline results weighted = model weighted.evaluate(test features, test labels,
                  batch size=128, verbose=0)
  for name, value in zip(model weighted.metrics names, baseline results weighted):
    print(name, ': ', value)
    print()
  plot cm(test labels, test predictions baseline weighted)
  return test predictions baseline weighted, baseline results weighted
test predictions baseline weighted 1, baseline results weighted 1 = nn(train df 1,
test df 1, train labels 1, train features 1, val features 1, val labels 1, test labels 1,
test features 1, initial bias 1, class weight 1)
test predictions baseline weighted 2, baseline results weighted 2 = nn(train df 2,
test df 2, train labels 2, train features 2, val features 2, val labels 2, test labels 2,
test features 2, initial bias 2, class weight 2)
test predictions baseline weighted 3, baseline results weighted 3 = nn(train df 3,
test df 3, train labels 3, train features 3, val features 3, val labels 3, test labels 3,
test features 3, initial bias 3, class weight 3)
upper bound(train df 1, train df labels 1, test df 1, test predictions baseline weighted 1)
#upper bound of nn for dataset a
upper bound(train df 2, train df labels 2, test df 2, test predictions baseline weighted 2)
#upper bound of nn for dataset b
upper bound(train df 3, train df labels 3, test df 3, test predictions baseline weighted 3)
#upper bound of nn for dataset c
```