

```
# Install Kaggle package
!pip install kaggle
# Make a directory for the Kaggle API key
!mkdir -p ~/.kaggle
# Move the Kaggle API key to the correct location
!cp kaggle.json ~/.kaggle/
# Set the permissions for the API key
!chmod 600 ~/.kaggle/kaggle.json
Requirement already satisfied: kaggle in
/usr/local/lib/python3.10/dist-packages (1.6.17)
Requirement already satisfied: six>=1.10 in
/usr/local/lib/python3.10/dist-packages (from kaggle) (1.16.0)
Requirement already satisfied: certifi>=2023.7.22 in
/usr/local/lib/python3.10/dist-packages (from kaggle) (2024.8.30)
Requirement already satisfied: python-dateutil in
/usr/local/lib/python3.10/dist-packages (from kaggle) (2.8.2)
Requirement already satisfied: requests in
/usr/local/lib/python3.10/dist-packages (from kaggle) (2.32.3)
Requirement already satisfied: tgdm in /usr/local/lib/python3.10/dist-
packages (from kaggle) (4.66.6)
Requirement already satisfied: python-slugify in
/usr/local/lib/python3.10/dist-packages (from kaggle) (8.0.4)
Requirement already satisfied: urllib3 in
/usr/local/lib/python3.10/dist-packages (from kaggle) (2.2.3)
Requirement already satisfied: bleach in
/usr/local/lib/python3.10/dist-packages (from kaggle) (6.2.0)
Requirement already satisfied: webencodings in
/usr/local/lib/python3.10/dist-packages (from bleach->kaggle) (0.5.1)
Requirement already satisfied: text-unidecode>=1.3 in
```

```
/usr/local/lib/python3.10/dist-packages (from python-slugify->kaggle)
(1.3)
Requirement already satisfied: charset-normalizer<4,>=2 in
/usr/local/lib/python3.10/dist-packages (from reguests->kaggle)
(3.4.0)
Requirement already satisfied: idna<4,>=2.5 in
/usr/local/lib/python3.10/dist-packages (from requests->kaggle) (3.10)
cp: cannot stat 'kaggle.json': No such file or directory
chmod: cannot access '/root/.kaggle/kaggle.json': No such file or
directory
# Download the dataset from Kaggle
!kaggle datasets download -d mlg-ulb/creditcardfraud
Dataset URL: https://www.kaggle.com/datasets/mlg-ulb/creditcardfraud
License(s): DbCL-1.0
Downloading creditcardfraud.zip to /content
97% 64.0M/66.0M [00:02<00:00, 34.2MB/s]
100% 66.0M/66.0M [00:02<00:00, 23.3MB/s]
# Unzip the downloaded file
!unzip creditcardfraud.zip
Archive: creditcardfraud.zip
replace creditcard.csv? [y]es, [n]o, [A]ll, [N]one, [r]ename:
import numpy as np
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
import matplotlib.gridspec as gridspec
from sklearn.ensemble import IsolationForest,RandomForestClassifier
from sklearn.model_selection import train test split,GridSearchCV
from sklearn.model selection import
KFold, cross val score, Stratified KFold
from sklearn.metrics import
accuracy score, precision score, recall score, f1 score
from sklearn.metrics import confusion matrix, ConfusionMatrixDisplay
from sklearn.metrics import
log_loss,roc_auc_score,f1_score,accuracy score
from sklearn.preprocessing import StandardScaler
from xgboost import XGBClassifier
from warnings import simplefilter
simplefilter("ignore")
# Load the dataset
data = pd.read csv('creditcard.csv')
```



```
# Displaying all the columns in the dataset
pd.set_option("display.max_columns",1000)

data.head()
{"type":"dataframe","variable_name":"data"}
```

Time--> column represents the time elapsed in seconds between each transaction and the first transaction in the dataset.

Columns V1-V28--> These are the result of a PCA dimensionality reduction. Their meaning has been made obscure intentionally because of privacy reasons.

Amount--> Transaction amount.

Class--> Type of transaction (1 for fraudulent, 0 for legit).

DATA EXPLORATION

```
data.info()
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 284807 entries, 0 to 284806
Data columns (total 31 columns):
     Column Non-Null Count
                              Dtype
             284807 non-null float64
     Time
             284807 non-null float64
 1
     ٧1
 2
    ٧2
             284807 non-null float64
3
    ٧3
             284807 non-null float64
    ۷4
             284807 non-null float64
    ۷5
             284807 non-null float64
 6
             284807 non-null float64
    ۷6
 7
    ٧7
             284807 non-null float64
 8
    ٧8
             284807 non-null float64
    ۷9
             284807 non-null float64
 10
    V10
             284807 non-null float64
             284807 non-null float64
 11
    V11
```

```
12
    V12
             284807 non-null float64
 13
    V13
             284807 non-null float64
 14 V14
             284807 non-null float64
             284807 non-null float64
 15
    V15
 16
    V16
             284807 non-null float64
17
    V17
             284807 non-null float64
18 V18
             284807 non-null float64
19 V19
             284807 non-null float64
 20 V20
             284807 non-null float64
21 V21
            284807 non-null float64
             284807 non-null float64
 22
    V22
23
   V23
             284807 non-null float64
 24 V24
             284807 non-null float64
 25
    V25
             284807 non-null float64
26 V26
             284807 non-null float64
             284807 non-null float64
 27
    V27
28 V28
            284807 non-null float64
            284807 non-null float64
29
    Amount
30 Class
            284807 non-null int64
dtypes: float64(30), int64(1)
memory usage: 67.4 MB
print("rows --->",data.shape[0],"no's")
print("columns -->",data.shape[1],"no's")
rows ---> 284807 no's
columns --> 31 no's
# Get the number of unique values for each column
unique counts = data.nunique()
print(unique counts)
          124592
Time
٧1
          275663
V2
          275663
V3
         275663
٧4
         275663
V5
          275663
۷6
          275663
٧7
          275663
V8
          275663
۷9
          275663
V10
          275663
V11
          275663
V12
         275663
V13
          275663
V14
          275663
          275663
V15
V16
          275663
V17
         275663
```

```
V18
          275663
V19
          275663
V20
          275663
V21
          275663
V22
          275663
V23
          275663
V24
          275663
V25
          275663
V26
          275663
V27
          275663
V28
          275663
Amount
          32767
Class
dtype: int64
data.describe()
# if cat columns then
# data.describe(include='object')
{"type": "dataframe"}
```

V1 to V28: These features have very small mean values (close to zero) and have been scaled and centered around zero. Their standard deviations range between 1 and 10, showing moderate variability.

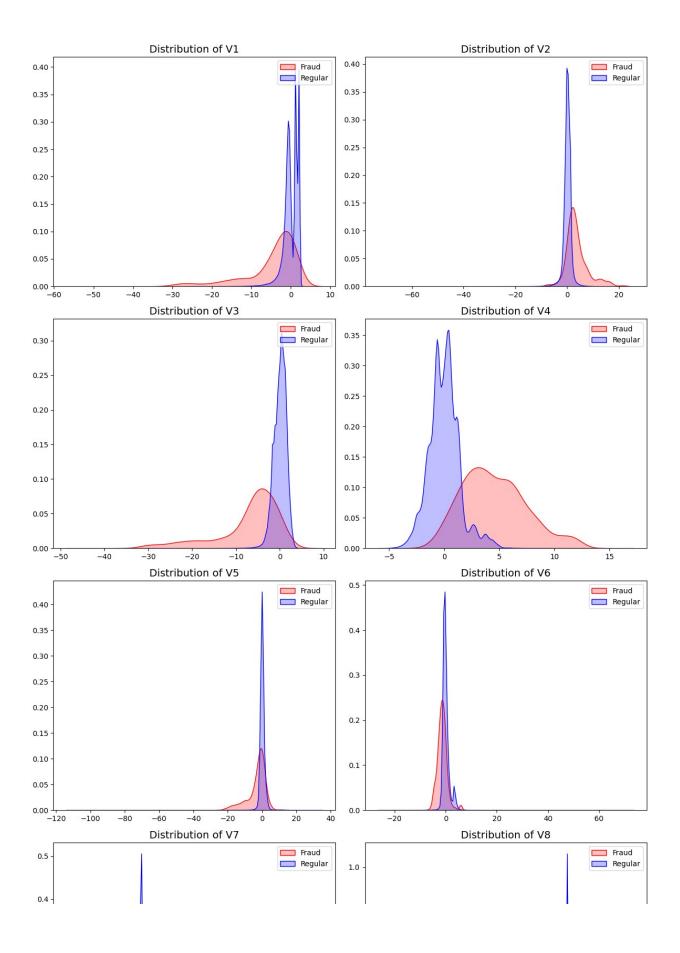
#Amount: Has a higher standard deviation than its mean, indicating a wide spread in transaction amounts (from very small to very large).

#Class: This categorical feature indicates whether a transaction is regular (0) or fraudulent (1). The mean of Class is close to zero because regular transactions are far more common than fraudulent ones

DISTRIBUTION OF V1... V28 Features

```
# Set up the plot grid for 28 features (V1 to V28) with 2 columns
num_features = 28
num_cols = 2
num_rows = num_features // num_cols + (num_features % num_cols > 0) #
Calculate number of rows needed
```

```
# Create the figure and axes
fig, axes = plt.subplots(num_rows, num_cols, figsize=(12, 5 *
num rows)) # Adjust height based on number of rows
# Flatten the axes array for easy indexing
axes = axes.flatten()
# Plot distributions for each of V1 to V28, separating by fraud and
regular transactions
for i in range(1, num features + 1):
    feature = f'V{i}'
      This loop iterates over the feature indices from 1 to 28.
# feature = f'V{i}': This creates a string representing the current
feature being plotted (e.g., 'V1', 'V2', ..., 'V28').
    # Plot for fraud transactions
    sns.kdeplot(data[data.Class == 1][feature], ax=axes[i - 1],
color='red', label='Fraud', fill=True)
    # Plot for regular transactions
    sns.kdeplot(data[data.Class == 0][feature], ax=axes[i - 1],
color='blue', label='Regular', fill=True)
    axes[i - 1].set title(f'Distribution of {feature}', fontsize=14)
    axes[i - 1].set xlabel('')
    axes[i - 1].set ylabel('')
    axes[i - 1].legend()
# Adjust layout to prevent overlap
plt.tight layout()
plt.show()
```



Overview of the Plots

X-axis: Each feature V 1 V1 to V 28 V28 will have its own plot, representing the values of that feature.

Y-axis: The density of the values, which shows how likely a particular value is to occur.

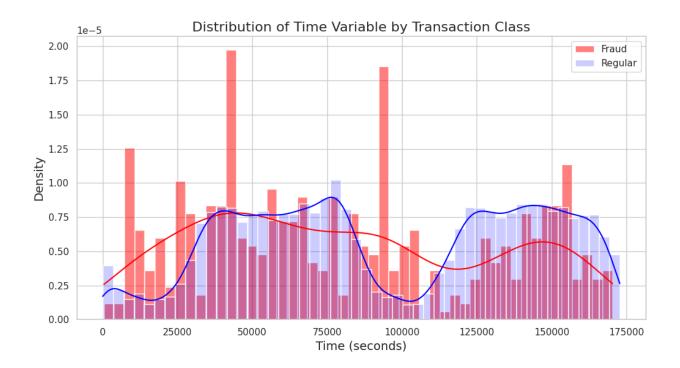
Classes:

Class 0 (Regular Transactions): This will be represented by one color (e.g., blue).

Class 1 (Fraudulent Transactions): This will be represented by another color (e.g., red).

DISTRIBUTION PLOT OF Time Variable

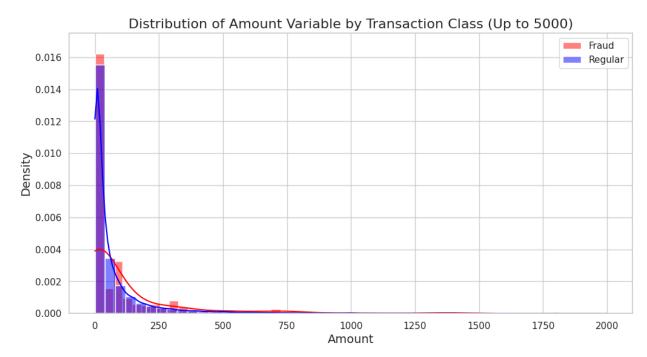
```
# Set the style for the plot
sns.set(style="whitegrid")
# Create a figure for the distribution plot
plt.figure(figsize=(12, 6))
# Plot the distribution of the 'Time' variable for fraud transactions
sns.histplot(data[data.Class == 1]['Time'], bins=50, kde=True,
color='red', stat='density', alpha=0.5, label='Fraud')
# Plot the distribution of the 'Time' variable for regular
transactions
sns.histplot(data[data.Class == 0]['Time'], bins=50, kde=True,
color='blue', stat='density', alpha=0.2, label='Regular')
# Set title and labels
plt.title('Distribution of Time Variable by Transaction Class',
fontsize=16)
plt.xlabel('Time (seconds)', fontsize=14)
plt.ylabel('Density', fontsize=14)
# Add legend
plt.legend()
# Show the plot
plt.show()
```



Distribution of Amount

```
# Set the style for the plot
sns.set(style="whitegrid")
# Create a figure for the distribution plot with increased height
plt.figure(figsize=(12, 6)) # Adjust height as needed
# Filter the data to include only amounts below or equal to 5000
amount filter = data[data['Amount'] <= 2000]</pre>
# Plot the distribution of the 'Amount' variable for fraud
transactions
sns.histplot(amount filter[amount_filter.Class == 1]['Amount'],
bins=50, kde=True, color='red', stat='density', alpha=0.5,
label='Fraud')
# Plot the distribution of the 'Amount' variable for regular
transactions
sns.histplot(amount filter[amount filter.Class == 0]['Amount'],
bins=50, kde=True, color='blue', stat='density', alpha=0.5,
label='Regular')
# Set title and labels
plt.title('Distribution of Amount Variable by Transaction Class (Up to
5000)', fontsize=16)
plt.xlabel('Amount', fontsize=14)
plt.ylabel('Density', fontsize=14)
```

```
# Adjust y-axis limits if needed
plt.ylim(0, 0.0175) # Adjust based on the data
# Add legend
plt.legend()
# Show the plot
plt.show()
```



Red (Fraud Transactions): This color is used to indicate transactions that are classified as fraudulent (where Class == 1). These transactions are typically of particular interest when analyzing credit card fraud data because they represent instances of fraudulent activity.

Blue (Regular Transactions): This color represents transactions that are classified as legitimate or regular (where Class == 0). These are the majority of transactions in the dataset and serve as a baseline for comparison against fraudulent transactions.

we visualized ->The Amount distribution for frauds is peaked on small quantities of money.



Duplicate Values

```
print('There are {} duplicate values in regular transactions out of
{}'.format(data[data['Class'] ==
0].duplicated().sum(),data[data['Class'] == 0].shape[0]))
print('There are {} duplicate values in fraudulent transactions out of
{}'.format(data[data['Class'] ==
1].duplicated().sum(),data[data['Class'] == 1].shape[0]))
There are 1062 duplicate values in regular transactions out of 284315
There are 19 duplicate values in fraudulent transactions out of 492
```

inplace=True: This modifies the DataFrame in place. The original DataFrame is updated, and no new DataFrame is returned. As a result, you do not need to assign the result to a new variable or even to the same variable.

inplace=False (default behavior): This will return a new DataFrame with the duplicates removed, leaving the original DataFrame unchanged. You would typically assign the result to a new variable or overwrite the existing variable.

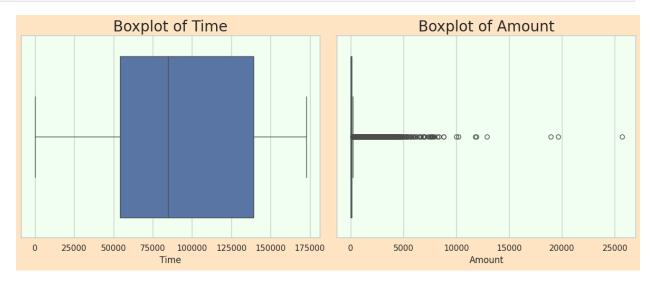
```
print('No. of rows before dropping duplicates: {}.'.format(len(data)))
data.drop_duplicates(inplace=True)
print('No. of rows after dropping duplicates: {}.'.format(len(data)))
No. of rows before dropping duplicates: 284807.
No. of rows after dropping duplicates: 283726.
```

here few duplicate rows dropped

HANDLING OUTLIERS

outlier can be extremely high or low values compared to the other observations and can be caused by measurement errors, natural variations in the data, or even unexpected discoveries.that leads lower performance so we deal with it...

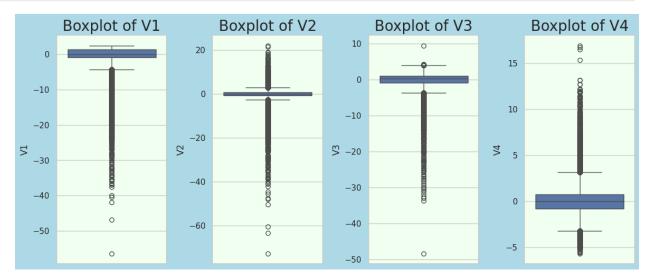
```
# Create subplots
fig, (ax1, ax2) = plt.subplots(ncols=2, figsize=(12, 5))
# Boxplot for Time
sns.boxplot(x=data['Time'], ax=ax1)
ax1.set title('Boxplot of Time', fontsize=20)
ax1.set facecolor('honeydew')
# Boxplot for Amount
sns.boxplot(x=data['Amount'], ax=ax2)
ax2.set title('Boxplot of Amount', fontsize=20)
ax2.set_facecolor('honeydew')
# Adjust layout
plt.tight layout()
# Set figure background color
fig.set facecolor('bisque')
# Show plot
plt.show()
```

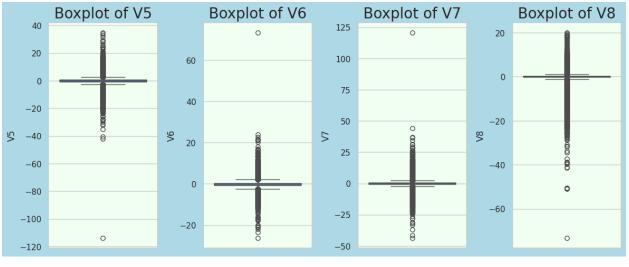


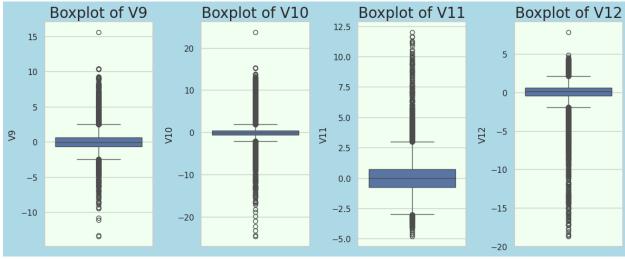
here no outlier in "time" but too many outlier in "amount"

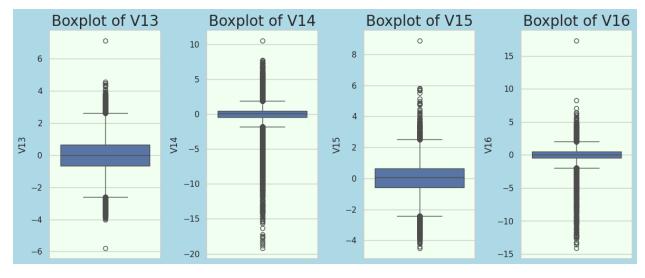
```
# Select only the V1-V28 features
features = list(data.columns.values)
del features[0]
```

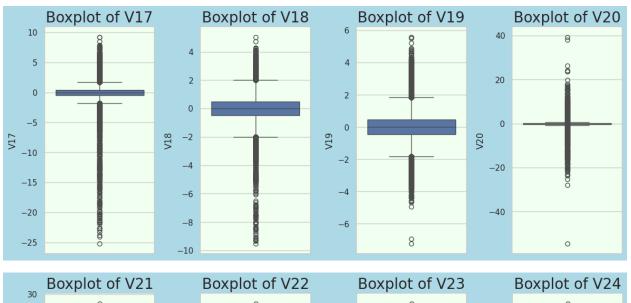
```
del features[28]
del features[28]
for i in range(7):#i \circ to \circ 6
  fig,
(ax1,ax2,ax3,ax4)=plt.subplots(ncols=4,figsize=(12,5))#wide=12,tall=5
  #plot diagram by 4 rows
  ax1=sns.boxplot(data[features[i*4]],ax=ax1)
  ax1.set_title('Boxplot of '+str(features[i*4]),fontsize=20)
  ax1.set facecolor('honeydew')
  ax2=sns.boxplot(data[features[i*4+1]],ax=ax2)
  ax2.set_title('Boxplot of '+str(features[i*4+1]),fontsize=20)
  ax2.set facecolor('honeydew')
 ax3=sns.boxplot(data[features[i*4+2]],ax=ax3)
  ax3.set_title('Boxplot of '+str(features[i*4+2]),fontsize=20)
  ax3.set facecolor('honeydew')
  ax4=sns.boxplot(data[features[i*4+3]],ax=ax4)
  ax4.set title('Boxplot of '+str(features[i*4+3]), fontsize=20)
  ax4.set facecolor('honeydew')
  plt.tight layout()
  fig.set facecolor('#ADD8E6')
```

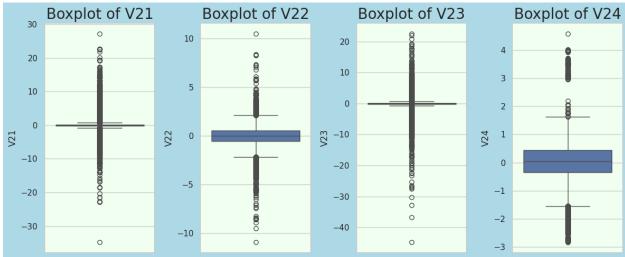


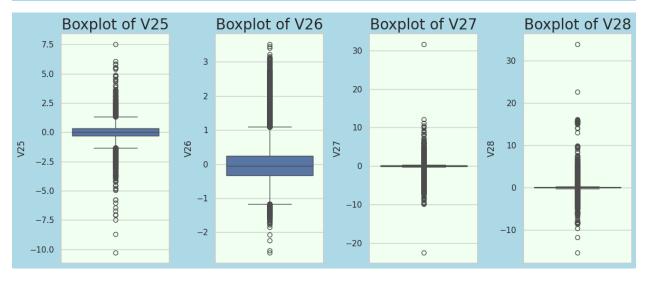












```
for column in features:
  #check how much skew
  print(f"Skewness for {column}: {data[column].skew()}")
Skewness for V1: -3.273271248440309
Skewness for V2: -4.6951619005404694
Skewness for V3: -2.1519839570997124
Skewness for V4: 0.6715041706728241
Skewness for V5: -2.414079246966253
Skewness for V6: 1.829880383771521
Skewness for V7: 2.890271192715498
Skewness for V8: -8.310970330052545
Skewness for V9: 0.5376630534496958
Skewness for V10: 1.2529670787468168
Skewness for V11: 0.34407419325686267
Skewness for V12: -2.1990082816149954
Skewness for V13: 0.06429340464018111
Skewness for V14: -1.9188037137586451
Skewness for V15: -0.3096590822936595
Skewness for V16: -1.0511614715174662
Skewness for V17: -3.690497194148406
Skewness for V18: -0.24866145737243997
Skewness for V19: 0.1083118109324772
Skewness for V20: -2.0431210560273323
Skewness for V21: 2.820033113572543
Skewness for V22: -0.18232972797521269
Skewness for V23: -5.867220791006341
Skewness for V24: -0.5521292366718961
Skewness for V25: -0.41574386205469593
Skewness for V26: 0.5802923172348093
Skewness for V27: -0.7538039138186547
Skewness for V28: 11.555115084196773
```

-ve indicates left skew

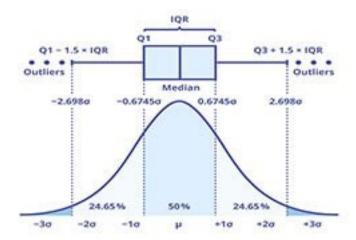
+ve indicates right skew

#CAPING OUTLIER

```
def outlier_imputer(data, features):
    data_out = data.copy()
    for column in features:
        # First define the first and third quartiles
        Q1 = data_out[column].quantile(0.25)
        Q3 = data_out[column].quantile(0.75)
        # Define the inter-quartile range
        IQR = Q3 - Q1
```

```
# ... and the lower/higher threshold values
        lowerL = Q1 - 1.5 * IQR
        higherL = Q3 + 1.5 * IQR
        # Impute 'left' outliers
        data out.loc[data out[column] < lowerL,column] = lowerL</pre>
        #which rows value lower than lowerL that capped
        # Impute 'right' outliers
        data_out.loc[data_out[column] > higherL,column] = higherL
        #data out.loc[mask, column] locates the rows in column where
data out[column] > higherL is True.-->[rows,columns]
    return data_out
#only made because of extracting features
data2 = data.drop('Class',axis=1)
feats = list(data2.columns.values) #list of column names
capped data = outlier imputer(data, feats)
pd.set option("display.max columns", 1000)
capped data.head()
{"type":"dataframe", "variable name": "capped data"}
```

the outliers have been imputed and the extreme values have been set to either Q 1 – 1.5 * I Q R (lower threshold) or to Q 3 + 1.5 * I Q R (higher threshold), where Q 1 and Q 3 are the first and third quartiles and I Q R is the so called interquartile range

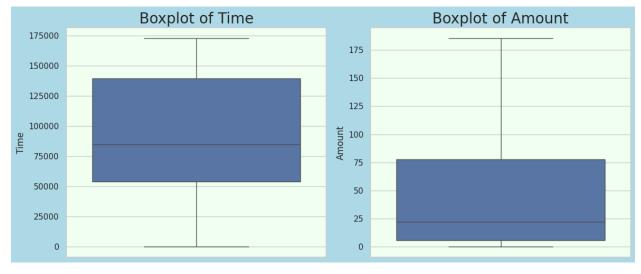


AFTER CAPPING

```
fig,(ax1,ax2) = plt.subplots(ncols=2,figsize=(12,5))
```

```
ax1 = sns.boxplot(capped_data['Time'],ax=ax1)
ax1.set_title('Boxplot of Time',fontsize=20)
ax1.set_facecolor('honeydew')
ax2 = sns.boxplot(capped_data['Amount'],ax=ax2)
ax2.set_title('Boxplot of Amount',fontsize=20)
ax2.set_facecolor('honeydew')

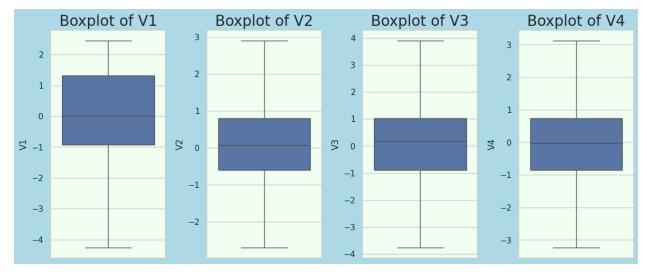
plt.tight_layout()
fig.set_facecolor('#ADD8E6')
```

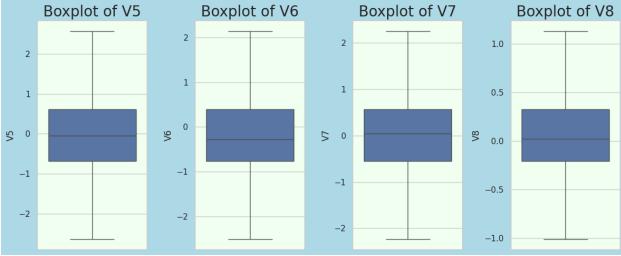


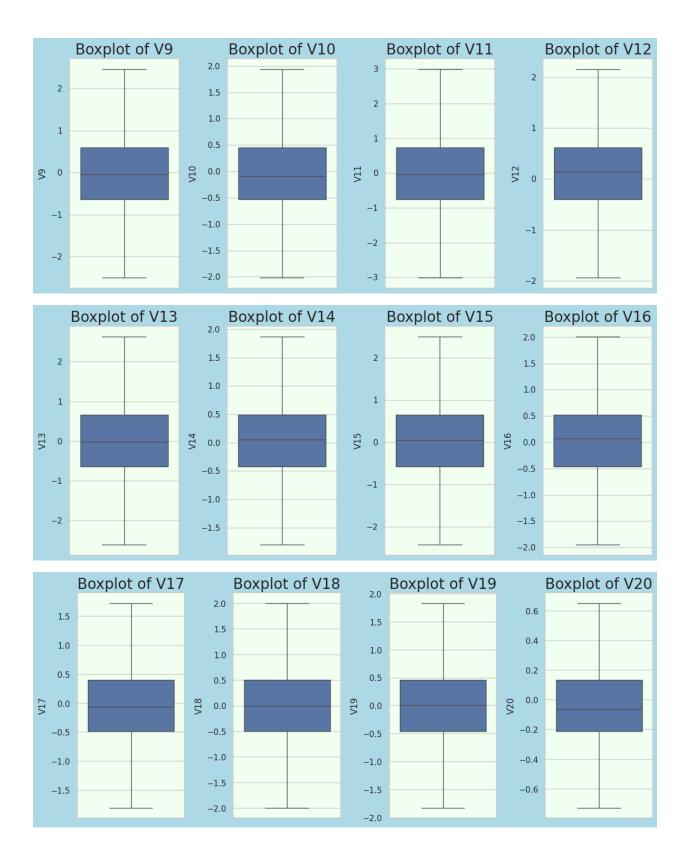
```
# Select only the V1-V28 features
features c = list(capped data.columns.values)
del features c[0]
del features c[28]
del features c[28]
for i in range(7):#i 0 to 6
(ax1,ax2,ax3,ax4)=plt.subplots(ncols=4,figsize=(12,5))#wide=12,tall=5
  #plot diagram by 4 rows
  ax1=sns.boxplot(capped data[features c[i*4]],ax=ax1)
  ax1.set_title('Boxplot of '+str(features_c[i*4]),fontsize=20)
  ax1.set facecolor('honeydew')
  ax2=sns.boxplot(capped data[features c[i*4+1]],ax=ax2)
  ax2.set title('Boxplot of '+str(features c[i*4+1]),fontsize=20)
  ax2.set facecolor('honeydew')
  ax3=sns.boxplot(capped data[features c[i*4+2]],ax=ax3)
  ax3.set_title('Boxplot of '+str(features_c[i*4+2]),fontsize=20)
  ax3.set facecolor('honeydew')
  ax4=sns.boxplot(capped data[features c[i*4+3]],ax=ax4)
```

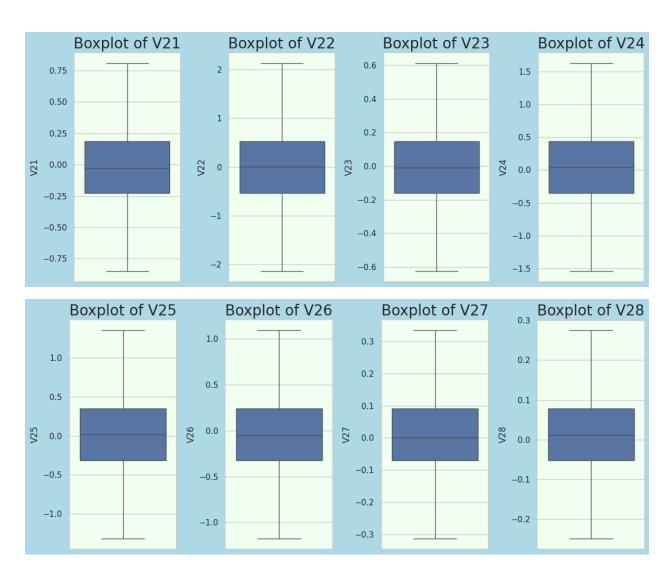
```
ax4.set_title('Boxplot of '+str(features_c[i*4+3]),fontsize=20)
ax4.set_facecolor('honeydew')

plt.tight_layout()
fig.set_facecolor('#ADD8E6')
```



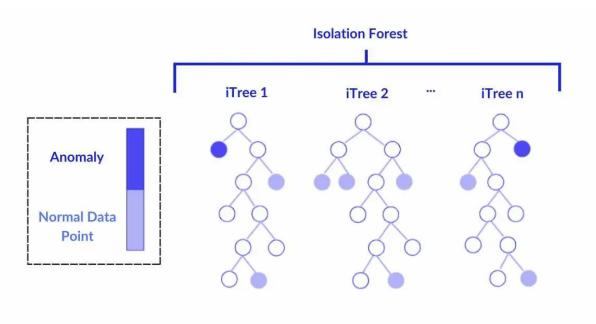






NOTE: after capping outliers, the data points are restricted to a defined range, which can potentially lead to a situation where many values are concentrated around the capped limits. This is where using anomaly detection techniques like Isolation Forest becomes especially important.

Isolation Forest & Dropping Outliers



```
data3 = data.copy()
data3 = data3.drop('Class',axis=1)

model =
IsolationForest(n_estimators=150,max_samples='auto',contamination=floa
t(0.1),max_features=1.0)
model.fit(data3)

IsolationForest(contamination=0.1, n_estimators=150)
```

Then, I am adding the score values and also an anomaly column to the dataframe. The negative score value of -1 indicates the presence of an anomaly. The value of 1 for the anomaly represents normal data.

You might set a threshold where scores less than 0 are considered anomalies, while scores equal to or greater than 0 are normal.

Calculating Scores: scores = model.decision_function(data3) computes the anomaly score for each data point in data3. The score reflects how isolated a data point is:

Points with lower scores are more likely to be anomalies.

scores that are more negative (e.g., -0.5, -1.0). These scores suggest that the point is more isolated, meaning it does not fit the typical pattern of the majority of the data.

```
anomaly = model.predict(data3)
anomaly
array([ 1,  1, -1, ...,  1,  1,  1])
```

generates binary predictions for the data points: The method returns -1 for points identified as anomalies and 1 for normal points.

```
#adding scores and anomalies
data3['score'] = scores
data3['anomaly'] = anomaly

data3.head()

{"type":"dataframe","variable_name":"data3"}

anomaly = data3.loc[data3['anomaly'] == -1]
print('The total number of outliers is {} out of
{}.'.format(len(anomaly),len(data)))

The total number of outliers is 28373 out of 283726.
```

dropping the outliers.

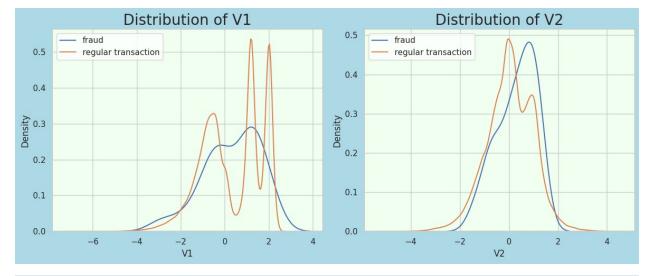
```
anomaly_index = list(anomaly.index)
forest_data = data.drop(anomaly_index,axis=0).reset_index(drop=True)
```

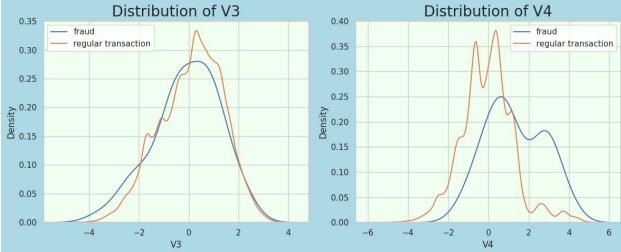
FINAL DISTRIBUTION

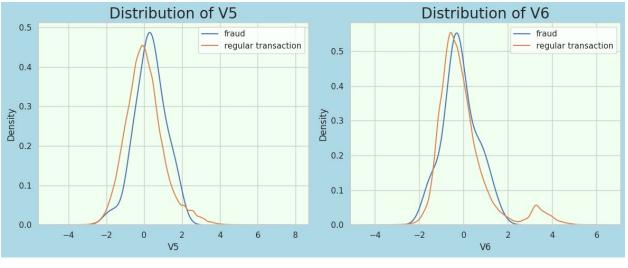
```
for i in range(14):
    fig,(ax1,ax2) = plt.subplots(ncols=2,figsize=(12,5))
    ax1 = sns.distplot(forest_data[forest_data.Class == 1]
[features[i*2]],ax=ax1,hist=False)
    ax1 = sns.distplot(forest_data[forest_data.Class == 0]
[features[i*2]],ax=ax1,hist=False)
    ax1.set_title('Distribution of '+str(features[i*2]),fontsize=20)
    ax1.set_facecolor('honeydew')
    ax1.legend(labels=['fraud','regular transaction'])
    ax2 = sns.distplot(forest_data[forest_data.Class == 1]
[features[i*2+1]],ax=ax2,hist=False)
    ax2 = sns.distplot(forest_data[forest_data.Class == 0]
```

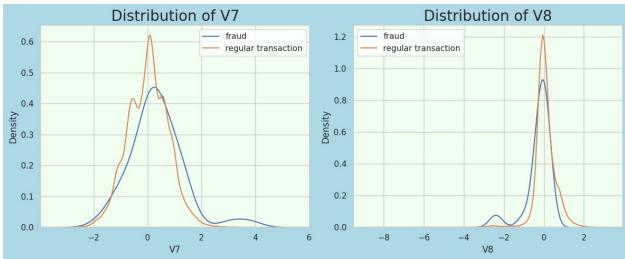
```
[features[i*2+1]],ax=ax2,hist=False)
   ax2.set_title('Distribution of '+str(features[i*2+1]),fontsize=20)
   ax2.set_facecolor('honeydew')
   ax2.legend(labels=['fraud','regular transaction'])

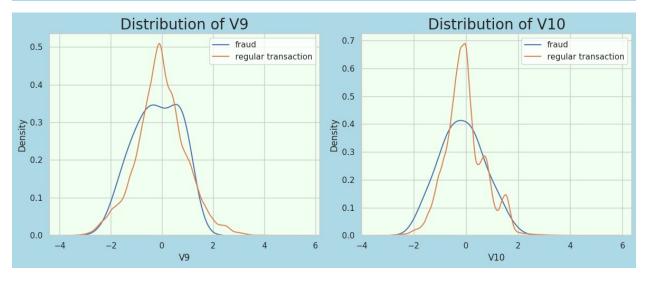
plt.tight_layout()
   fig.set_facecolor('#ADD8E6')
```

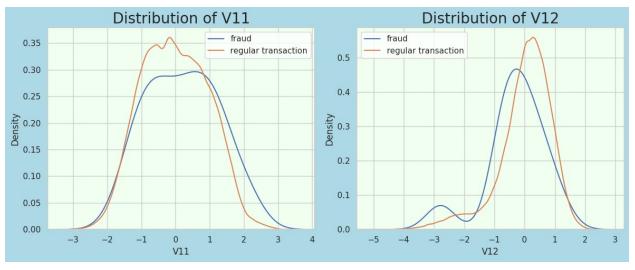


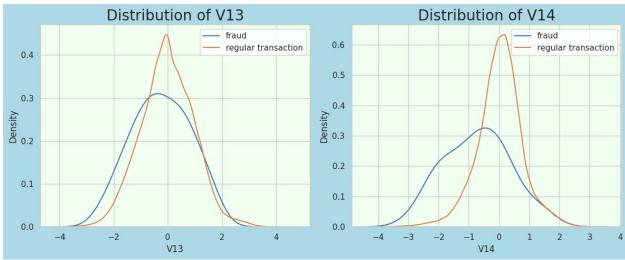


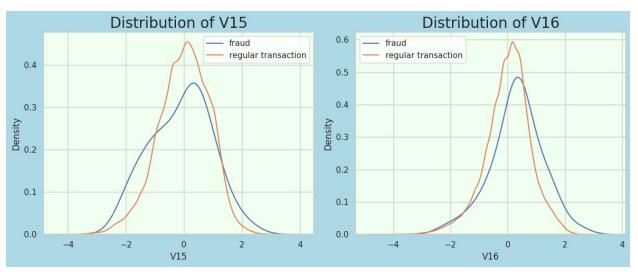


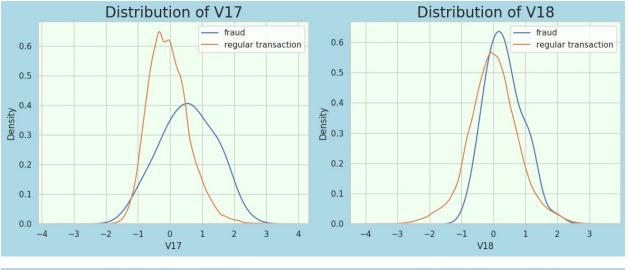


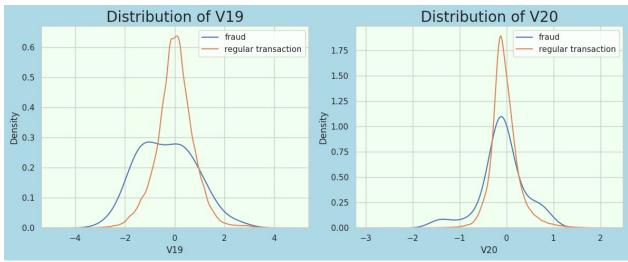


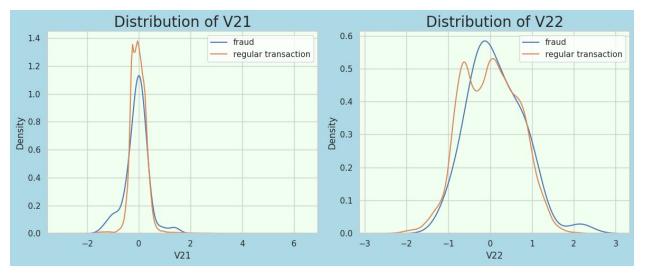


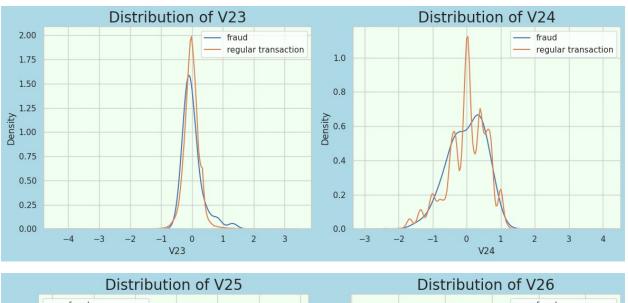


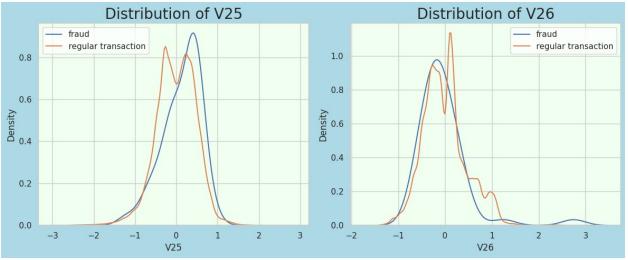


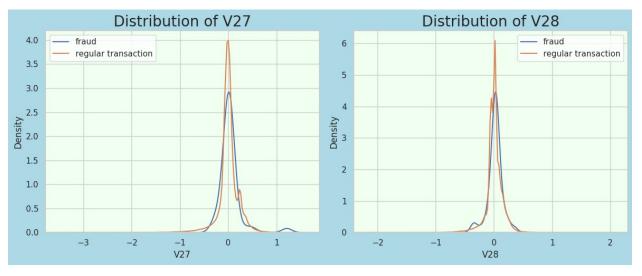




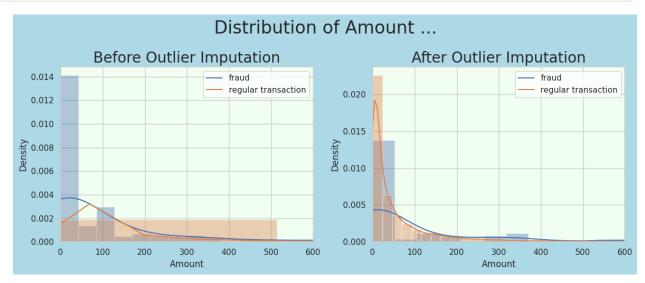








```
fig,(ax1,ax2) = plt.subplots(ncols=2,figsize=(12,5))
#BEFORE
ax1 = sns.distplot(data['Amount'][data.Class == 1],ax=ax1)
ax1 = sns.distplot(data['Amount'][data.Class == 0],ax=ax1)
ax1.set xlim(0,600)
ax1.set_title('Before Outlier Imputation',fontsize=20)
ax1.legend(labels=['fraud','regular transaction'])
ax1.set facecolor('honeydew')
#AFTER
ax2 = sns.distplot(forest data['Amount'][forest data.Class ==
1],ax=ax2)
ax2 = sns.distplot(forest data['Amount'][forest data.Class ==
[0],ax=ax2)
ax2.set xlim(0,600)
ax2.set title('After Outlier Imputation', fontsize=20)
ax2.legend(labels=['fraud','regular transaction'])
ax2.set facecolor('honeydew')
#PLOT
fig.suptitle("Distribution of Amount ...", fontsize=24)
plt.tight layout()
fig.set facecolor('#ADD8E6')
```



Before Outlier Imputation:

In the left plot, some transactions have very high amounts, which makes the data look stretched out to the right. Regular transactions usually have smaller amounts, while fraud transactions sometimes go higher.

After Outlier Imputation:

In the right plot, those high-amount transactions have been reduced or adjusted, so the data is now focused more on smaller amounts. This makes the graph more compact, with fewer large transactions.

In short: MACHINE FOCOUS ON HIGH VALUE SO,

We reduced the impact of big, unusual transaction amounts, so now the data focuses more on typical, smaller amounts. This can help make patterns in the data clearer.

FEATURE IMPORTANCE

identify which features are most influential in making predictions.

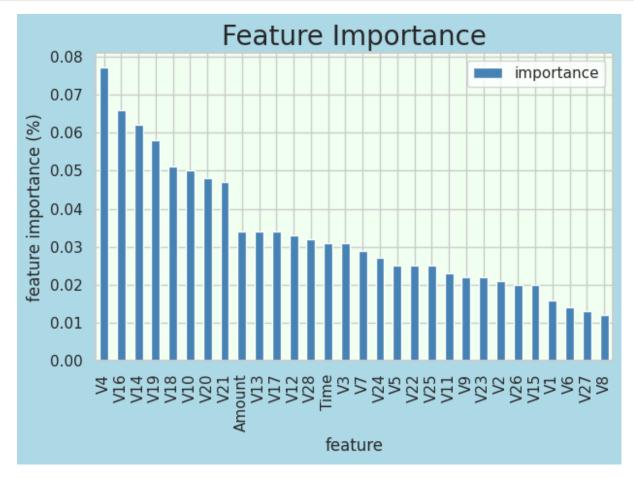
```
X = forest_data.drop('Class',axis=1)
y = forest_data['Class']
```

I can look at the relative importance of the features by means of a random forest classifier.

```
#USING RANDOM FOREST FIND FEATURE IMPORTANCE
# Random Forest Model
random forest = RandomForestClassifier(random state=1, max depth=4)
random forest.fit(X,y)
RandomForestClassifier(max depth=4, random state=1)
FI=random forest.feature importances
importances =
pd.DataFrame({'feature':X.columns,'importance':np.round(FI,3)})
importances =
importances.sort values('importance',ascending=False).set index('featu
importances.head(30)
{"summary":"{\n \"name\": \"importances\",\n \"rows\": 30,\n
\"fields\": [\n {\n \"column\": \"feature\",
\"properties\": {\n \"dtype\": \"string\",\n
                            \"column\": \"feature\",\n
\"num unique values\": 30,\n
                                      \"samples\": [\n
                                                                 \"V6\",\
           \<u>"</u>V7\",\n
                                \"V2\"\n
n \"V7\",\n \"V2\"\n ],\n
\"semantic_type\": \"\",\n \"description\": \"\"\n
0.077,\n \"num_unique_values\": 23,\n \"samp 0.023,\n 0.033,\n 0.077\n ],\n \"semantic_type\": \"\",\n \"description\": \"\"\n
     }\n ]\n}","type":"dataframe","variable_name":"importances"}
importances.plot.bar(color='steelblue')
```

```
plt.ylabel('feature importance (%)',fontsize=12)
plt.title('Feature Importance',fontsize=20)

plt.tight_layout()
plt.gcf().patch.set_facecolor('#ADD8E6')
plt.gca().set_facecolor('honeydew')
plt.show()
```



TO AVOID OVERFITTING: Apparently, there are three dominant features: V14, V17 and V4. All the others are much less important. SO WE DROP SOME COLUMNS

#experiment:

without drop any column random_forest accuracy --0.9997 DRP -'V1','V2','V3','V5','V6','V7','V8','V9','V11','V13','V14','V15','V16','V18','V19','V20','V21', 'V22','V23','V24','V25','V26','V27','V28' random_forest accuracy --->0.9997

so no DIFFERENCE

```
# Train-test split
X train,X test,y train,y test =
train_test_split(X,y,test_size=0.3,random_state=0)
random forest = RandomForestClassifier(class weight='balanced')
random forest.fit(X train,y train)
RandomForestClassifier(class weight='balanced')
def get test scores(model name:str,preds,y test data):
   accuracy = accuracy score(y test data,preds)
   precision = precision score(y test data,preds,average='macro')
   recall
             = recall_score(y_test_data,preds,average='macro')
             = f1 score(y test data,preds,average='macro')
   table = pd.DataFrame({'model': [model name], 'precision':
[precision], 'recall': [recall],
                        'F1': [f1], 'accuracy': [accuracy]})
    return table
# Use the model to predict on test data
rf test preds = random forest.predict(X test)
rf test results = get test scores('RF (test)',rf test preds,y test)
rf test results
{"summary":"{\n \"name\": \"rf test results\",\n \"rows\": 1,\n
\"fields\": [\n {\n \"column\": \"model\",\n
\"properties\": {\n \"dtype\": \"string\",\n
\"RF
},\n {\n \"column\":
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\"num_unique_values\": 1,\n
\"samples\": [\n
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\"recall\",\n \"properties\": {\n \"std\": null,\n \"min\": 0.5,\n
                                         \"dtype\": \"number\",\n
\"max\": 0.5,\n
\"num_unique_values\": 1,\n \"samples\": [\n
                                                         0.5\n
           \"semantic_type\": \"\",\n \"description\": \"\"\n
],\n
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}\n },\n {\n
\"dtype\": \"number\",\n \"std\": null,\n \"m: 0.4999477793153868,\n \"max\": 0.4999477793153868,\n
                                                     \"min\":
\"num_unique_values\": 1,\n \"samples\": [\n
0.4999477793153868\n ],\n
                                     \"semantic type\": \"\",\n
\"description\": \"\"\n
                                         {\n \"column\":
                          }\n
                                 },\n
```

```
\"accuracy\",\n \"properties\": {\n \"dtype\":
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\"num_unique_values\": 1,\n \"samples\": [\n
0.9997911390752683\n ],\n \"semantic_type\": \"\",\n
\"description\": \"\"\n }\n }\n ]\
n}","type":"dataframe","variable_name":"rf_test_results"}
```

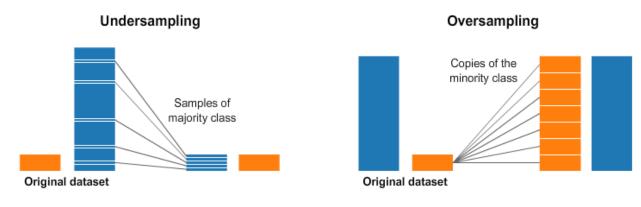
#Accuracy gives a general idea of how well the model performs overall.

#Precision is important when the cost of false positives is high (e.g., in spam detection).

#Recall is crucial when the cost of false negatives is high (e.g., in disease detection).

#F1 Score is a balanced measure that is useful when you need a balance between precision and recall, especially when the class distribution is imbalanced.

Resampling and Cross Validation



Undersampling

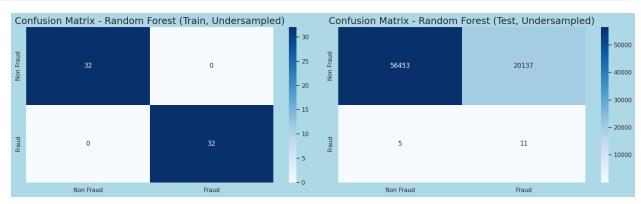
```
from imblearn.under_sampling import RandomUnderSampler
# Create a RandomUnderSampler object
rus = RandomUnderSampler(random_state=42,sampling_strategy='majority')
# Balancing the data
X_resampled,y_resampled = rus.fit_resample(X_train,y_train)# sample we used
```

Random Forest Classifier

```
random_forest.fit(X_resampled,y_resampled)
# Use the model to predict on train data
rf_train_resampled_preds = random_forest.predict(X_resampled)
```

```
# Use the model to predict on test data
rf test preds = random forest.predict(X test)
rf test results = get test scores('RF (test)',rf_test_preds,y_test)
rf test results
{"summary":"{\n \"name\": \"rf_test_results\",\n \"rows\": 1,\n
                        \column\": \mbox{"model}",\n
\"fields\": [\n {\n
                        \"dtype\": \"string\",\n
\"properties\": {\n
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                                                         \"RF
                                \"samples\": [\n
                           \"semantic_type\": \"\",\n
             ],\n
(test)\"\n
\"description\": \"\"\n }\n
                                },\n {\n \"column\":
                                        \"dtype\":
\"precision\",\n \"properties\": {\n
                  \"std\": null,\n \"min\":
\"number\",\n
\"num_unique_values\": 1,\n \"samples\": [\n
0.5002286992441416\n ],\n
                                \"semantic_type\": \"\",\n
                               },\n {\n \"column\":
\"description\": \"\"\n
                          }\n
\"recall\",\n \"properties\": {\n \"dtype\": \"number\",\n \"std\": null,\n \"min\": 0.7122902794098447,\n \"max\":
                                          \"dtype\": \"number\",\n
\"num unique values\": 1,\n
\"semantic_type\": \"\",\n \"description\": \"\"\n
    \"dtype\": \"number\",\n \"std\": null,\n \"min\\0.42485104048074507,\n \"max\": 0.42485104048074507,\n \"num_unique_values\": 1,\n \"samples\": [\n
                                                   \"min\":
                          ],\n \"semantic_type\": \"\",\n
}\n },\n {\n \"column\":
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\"description\": \"\"\n
                                 },\n {\n
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\"number\",\n \"std\": null,\n
                                         \"dtype\":
                                     \"min\":
0.7370702033783254,\n\\"max\": 0.7370702033783254,\n
\"num_unique_values\": 1,\n
                                \"samples\": [\n
0.7370702033783254\n
                                     \"semantic_type\": \"\",\n
                          ],\n
\"description\": \"\"\n
                          n}","type":"dataframe","variable_name":"rf_test_results"}
fig, (ax1, ax2) = plt.subplots(ncols=2, figsize=(17, 5))
# Generate array of values for confusion matrix
cm1 =
confusion matrix(y resampled,rf train resampled preds,labels=random fo
rest.classes )
# Plot confusion matrix for the training set
sns.heatmap(cm1, annot=True, ax=ax1, fmt='d', cmap='Blues')
ax1.xaxis.set_ticklabels(['Non Fraud', 'Fraud'])
ax1.yaxis.set_ticklabels(['Non Fraud', 'Fraud'])
ax1.set_title('Confusion Matrix - Random Forest (Train,
Undersampled)', fontsize=18)
ax1.set facecolor('honeydew')
```

```
# Generate array of values for confusion matrix
cm2 =
confusion_matrix(y_test,rf_test_preds,labels=random_forest.classes_)
# Plot confusion matrix for the test set
sns.heatmap(cm2, annot=True, ax=ax2, fmt='d', cmap='Blues')
ax2.xaxis.set_ticklabels(['Non Fraud', 'Fraud'])
ax2.yaxis.set ticklabels(['Non Fraud', 'Fraud'])
ax2.set title('Confusion Matrix - Random Forest (Test, Undersampled)',
fontsize=18)
ax2.set facecolor('honeydew')
#plot
# Set figure background color
plt.gcf().patch.set facecolor('#ADD8E6')
# Adjust layout
plt.tight layout()
plt.show()
```

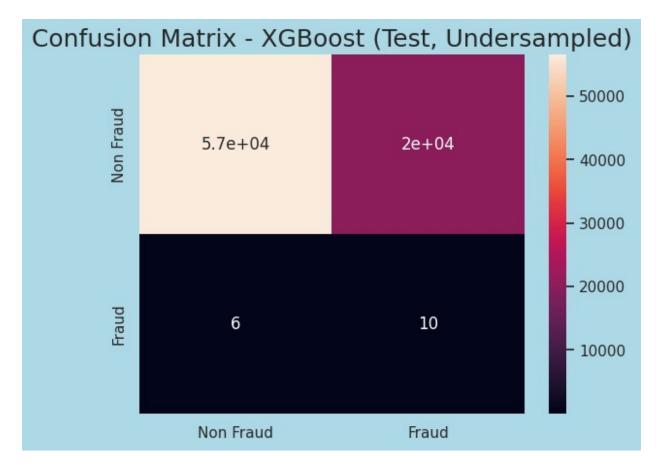


XGBoost Classifier

```
# Instantiate the XGBoost classifier
xgb = XGBClassifier(objective='binary:logistic',random_state=42)
xgb.fit(X_resampled,y_resampled)
# Use the model to predict on train data
xgb_train_resampled_preds = xgb.predict(X_resampled)
# Use the model to predict on test data
xgb_test_preds = xgb.predict(X_test)

xgb_test_results = get_test_scores('XGB (test)',xgb_test_preds,y_test)
xgb_test_results
{"summary":"{\n \"name\": \"xgb_test_results\",\n \"rows\": 1,\n
\"fields\": [\n {\n \"column\": \"model\",\n
\"properties\": {\n \"dtype\": \"string\",\n
```

```
\"XGB
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\"samples\": [\n
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\"num_unique_values\": 1,\n \"samples\": [\n
n}","type":"dataframe","variable_name":"xgb_test_results"}
# Generate array of values for confusion matrix
cm = confusion_matrix(y_test,xgb test preds,labels=xgb.classes )
ax = sns.heatmap(cm,annot=True)
ax.xaxis.set ticklabels(['Non Fraud','Fraud'])
ax.yaxis.set ticklabels(['Non Fraud','Fraud'])
ax.set title('Confusion Matrix - XGBoost (Test,
Undersampled)',fontsize=18)
plt.gcf().patch.set facecolor('#ADD8E6')
```



Oversampling

```
from imblearn.over_sampling import SMOTE

smote = SMOTE()

# Balancing the data
X_oversampled,y_oversampled = smote.fit_resample(X_train,y_train)
```

Random Forest *Classifier*

```
random_forest.fit(X_oversampled,y_oversampled)

# Use the model to predict on train data
rf_train_oversampled_preds = random_forest.predict(X_oversampled)

rf_train_oversampled_results = get_test_scores('RF (train,
oversampled)',rf_train_oversampled_preds,y_oversampled)

rf_train_oversampled_results

{"summary":"{\n \"name\": \"rf_train_oversampled_results\",\n
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\"properties\": {\n \"dtype\": \"string\",\n
\"num_unique_values\": 1,\n \"samples\": [\n \"RF
```

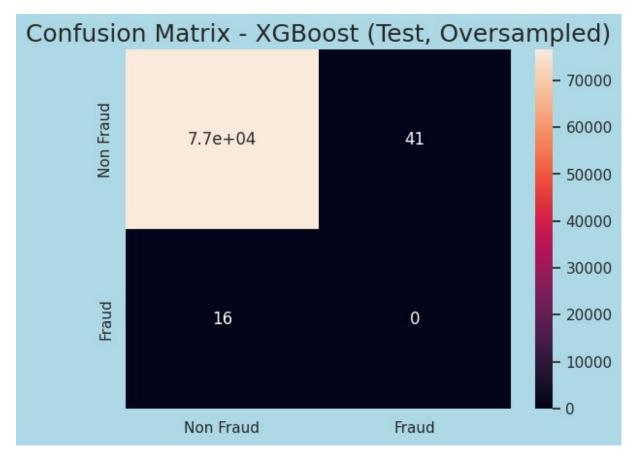
```
(train, oversampled)\"\n ],\n
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                                                                                                                  {\n \"column\":
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\"description\": \"\"\n }\n
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\"num_unique_values\": 1,\n \"samples\": [\n 1.0\n
],\n \"semantic_type\": \"\",\n \"description\": \"\"\n
}\n    },\n    {\n     \"column\": \"F1\",\n     \"properties\": {\n
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n     \"max\": 1.0,\n     \"num_unique_values\": 1,\n
\''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \''', \ \'''
\"column\": \"accuracy\",\n \"properties\": {\n \"dtype\":
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n}","type":"dataframe","variable name":"rf train oversampled results"}
# Use the model to predict on test data
rf test preds = random forest.predict(X test)
rf test results = get test scores('RF (test)', rf test preds, y test)
rf test results
{"summary":"{\n \"name\": \"rf_test_results\",\n \"rows\": 1,\n
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                                                                                                                                                              \"RF
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```

```
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                                          \"semantic type\": \"\",\n
                                     },\n {\n \"column\":
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                                         \"semantic type\": \"\",\n
                                     }\n ]\
n}","type":"dataframe","variable name":"rf test results"}
```

XGBoost Classifier

```
# 1. Instantiate the XGBoost classifier
xqb = XGBClassifier(objective='binary:logistic',random state=42)
xgb.fit(X oversampled,y oversampled)
XGBClassifier(base score=None, booster=None, callbacks=None,
              colsample bylevel=None, colsample bynode=None,
              colsample bytree=None, device=None,
early stopping rounds=None,
              enable categorical=False, eval metric=None,
feature types=None,
              gamma=None, grow_policy=None, importance_type=None,
              interaction constraints=None, learning rate=None,
max_bin=None,
              max cat threshold=None, max cat to onehot=None,
             max delta step=None, max depth=None, max leaves=None,
             min child weight=None, missing=nan,
monotone constraints=None,
              multi strategy=None, n estimators=None, n jobs=None,
              num parallel tree=None, random state=42, ...)
# Use the model to predict on test data
xgb test preds = xgb.predict(X test)
xgb test results = get test_scores('XGB (test)',xgb_test_preds,y_test)
xgb test results
{"summary":"{\n \"name\": \"xgb_test_results\",\n \"rows\": 1,\n
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(test)\"\n
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                     ],\n
                                        \"semantic type\": \"\",\n
```

```
\"description\": \"\"\n
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\"num unique values\": 1,\n \"samples\": [\n
0.999\overline{2559329\overline{556432\n} \"description\":\"\"n \\n\\"se
                                 \"semantic type\": \"\",\n
n}","type":"dataframe","variable name":"xgb test results"}
# Generate array of values for confusion matrix
cm = confusion matrix(y test,xgb test preds,labels=xgb.classes )
ax = sns.heatmap(cm,annot=True)
ax.xaxis.set ticklabels(['Non Fraud', 'Fraud'])
ax.yaxis.set ticklabels(['Non Fraud','Fraud'])
ax.set title('Confusion Matrix - XGBoost (Test,
Oversampled)',fontsize=18)
plt.gcf().patch.set facecolor('#ADD8E6')
```



It seems that undersampling improves the classification of the minority class (frauds), while oversampling tends to do better on the majority class (non frauds, i.e. regular transactions).

In light of this, I will keep the undersampling strategy and try to improve its results by also implementing other strategies.

since undersampling random forest give best recall score so we cross validate to this and check score

```
# K-fold stratified cross validation
kfold = StratifiedKFold(n_splits=10)
kfold

StratifiedKFold(n_splits=10, random_state=None, shuffle=False)

#GRID SEARCH CV TAKE TOO MUCH TIM SO I COMMENT OUT IF U HAVE 6 CORE
PROCESSOR THEN IT WILL BE DONE IN 20 MIN. ACCURACY IS NEAR TO -->0.77

# RFC Parameters tuning
# RFC = RandomForestClassifier(random_state=42)

# rf_param_grid = {
        'max_depth': [2,3,4,5,None],
        'max_features': [1.0],
        'max_samples': [1.0],
```

```
'min samples leaf': [2,3,4],
      'min samples split': [2,2,4],
#
#
      'n_estimators': [200,300,400]
# qsRFC =
GridSearchCV(RFC, param_grid=rf_param_grid, cv=kfold, scoring="f1")
# gsRFC.fit(X resampled, y resampled)
#FOR TIME TAKING I USE RANDOM SEARCH CV
from sklearn.model selection import RandomizedSearchCV
RFC = RandomForestClassifier(random state=42)
# Define the Stratified K-Folds cross-validator
kfold = StratifiedKFold(n splits=10)
# Define the parameter grid for RandomizedSearchCV
rf param dist = {
    'max_depth': [2, 3, 4, 5, None],
    'max_features': [1.0],
    'max samples': [1.0],
    'min samples leaf': [2, 3, 4],
    'min samples split': [2, 4],
    'n_estimators': [200, 300, 400]
}
# Initialize RandomizedSearchCV
rsRFC = RandomizedSearchCV(RFC, param_distributions=rf_param_dist,
n iter=100, cv=kfold, scoring="f1", random state=42)
# Fit the model
rsRFC.fit(X resampled, y resampled)
# Optionally, you can retrieve the best parameters
best params = rsRFC.best params
print("Best parameters found: ", best params)
# Use the model to predict on test data
rf test preds = rsRFC.predict(X test)
rf test results = get test scores('RF (test)', rf test preds, y test)
rf test results
# Generate array of values for confusion matrix
cm = confusion matrix(y test,rf test preds,labels=rsRFC.classes )
ax = sns.heatmap(cm,annot=True)
ax.xaxis.set_ticklabels(['Non Fraud','Fraud'])
ax.yaxis.set_ticklabels(['Non Fraud','Fraud'])
ax.set title('Confusion Matrix - Random Forest (Test, Undersampled &
```

```
CV)',fontsize=18)
plt.gcf().patch.set_facecolor('#ADD8E6')
```

Till now Best Recall Score is --> under sampling, random forest --> recall score of 0.788003 for your credit card fraud detection model using Random Forest with undersampling! A high recall is crucial in fraud detection, as it indicates that your model is effectively identifying fraudulent transactions.

```
# !pip install catboost
# from sklearn.ensemble import VotingClassifier
# from sklearn.ensemble import RandomForestClassifier
# from lightgbm import LGBMClassifier
# from catboost import CatBoostClassifier
# from xgboost import XGBClassifier
# # Instantiate individual classifiers
# rf = RandomForestClassifier(n estimators=100, random state=42)
# lgbm = LGBMClassifier(objective='binary', random state=42)
# xgb = XGBClassifier(objective='binary:logistic', random state=42)
# catboost = CatBoostClassifier(iterations=1000, learning rate=0.1,
depth=6, random seed=42, silent=True)
# # Instantiate the Voting Classifier
# voting clf = VotingClassifier(
     estimators=[
          ('rf', rf),
          ('lgbm', lgbm),
          ('xgb', xgb),
          ('catboost', catboost)
     voting='hard' # Use 'soft' for soft voting
# # Fit the Voting Classifier on the resampled training data
# voting_clf.fit(X_resampled, y_resampled)
# # Use the model to predict on train data
```

```
# voting_train_resampled_preds = voting_clf.predict(X_resampled)
# # Use the model to predict on test data
# voting_test_preds = voting_clf.predict(X_test)
# # Get test results using your scoring function
# voting_test_results = get_test_scores('Voting Classifier (test)',
voting_test_preds, y_test)
# voting_test_results
```