## **Credit Card Fraud Detection** ¶

Using a dataset of of nearly 28,500 credit card transactions and multiple unsupervised anomaly detection algorithms, we are going to identify transactions with a high probability of being credit card fraud. In this project, we shall build and deploy the following two machine learning algorithms:

- 1. Local Outlier Factor (LOF)
- 2. Isolation Forest Algorithm

Furthermore, we shall use metrics suchs as precision, recall, and F1-scores.

In addition, we shall explore parameter histograms and correlation matrices.

```
In [1]: #importing necessary libraries
        import sys
        import numpy
        import pandas
        import matplotlib
        import seaborn
        import scipy
        print('Python: {}'.format(sys.version))
        print('Numpy: {}'.format(numpy. version ))
        print('Pandas: {}'.format(pandas. version ))
        print('Matplotlib: {}'.format(matplotlib. version ))
        print('Seaborn: {}'.format(seaborn. version ))
        print('Scipy: {}'.format(scipy. version ))
        Python: 3.7.0 (default, Jun 28 2018, 08:04:48) [MSC v.1912 64 bit (AMD64)]
        Numpy: 1.16.4
        Pandas: 0.23.4
        Matplotlib: 2.2.3
        Seaborn: 0.9.0
        Scipy: 1.2.1
```

```
In [2]: # import the necessary packages
        import numpy as np
        import pandas as pd
        import matplotlib.pyplot as plt
        import seaborn as sns
In [3]: # Load the dataset from the csv file using pandas
        data = pd.read csv('creditcard.csv')
In [4]: # Start exploring the dataset
        print(data.columns)
        print(data.shape)
        Index(['Time', 'V1', 'V2', 'V3', 'V4', 'V5', 'V6', 'V7', 'V8', 'V9', 'V10',
               'V11', 'V12', 'V13', 'V14', 'V15', 'V16', 'V17', 'V18', 'V19', 'V20',
               'V21', 'V22', 'V23', 'V24', 'V25', 'V26', 'V27', 'V28', 'Amount',
               'Class'],
              dtype='object')
        (284807, 31)
```

```
In [5]: # Print the shape of the data
data = data.sample(frac=0.1, random_state = 1) #since original dataset is too large
print(data.shape)
print(data.describe())

# V1 - V28 are the results of a PCA Dimensionality reduction to protect user identities and sensitive features
```

(28481, 31)V1 V2 V4 \ Time V3 28481.000000 count 28481.000000 28481.000000 28481.000000 28481.000000 94705.035216 -0.001143 -0.018290 0.000795 0.000350 mean std 47584.727034 1.994661 1.709050 1.522313 1.420003 min 0.000000 -40.470142 -63,344698 -31.813586 -5.266509 25% 53924.000000 -0.908809 -0.610322 -0.892884 -0.847370 50% 84551.000000 0.031139 0.051775 0.178943 -0.017692 75% 139392,000000 1.320048 0.792685 1.035197 0.737312 172784.000000 2.411499 17.418649 4.069865 16.715537 max V5 V6 V7 V8 V9 \ count 28481.000000 28481.000000 28481.000000 28481.000000 28481,000000 -0.015666 0.003634 -0.008523 -0.003040 0.014536 mean std 1.395552 1.334985 1.237249 1.204102 1.098006 -42.147898 -19.996349 -22.291962 -8.739670 min -33.785407 25% -0.703986 -0.765807 -0.562033 -0.208445 -0.632488 50% -0.068037 -0.269071 0.028378 0.024696 -0.037100 75% 0.603574 0.398839 0.559428 0.326057 0.621093 28.762671 22.529298 36.677268 19.587773 8.141560 max V21 V22 V23 V24 \ . . . 28481,000000 count 28481.000000 28481.000000 28481.000000 . . . 0.004740 0.006719 -0.000494 -0.002626 mean . . . std 0.744743 0.728209 0.645945 0.603968 min -16.640785 -10.933144 -30.269720 -2.752263 25% -0.224842 -0.535877 -0.163047 -0.360582 50% -0.029075 0.014337 -0.012678 0.038383 . . . 75% 0.533936 0.434851 0.189068 0.148065 . . . 22.588989 6.090514 15.626067 3.944520 max . . . V25 V26 V27 V28 Amount \ count 28481.000000 28481.000000 28481.000000 28481.000000 28481.000000 -0.000917 0.004762 -0.001689 -0.004154 89.957884 mean std 0.520679 0.488171 0.418304 0.321646 270.894630 -2.534330 0.000000 min -7.025783 -8.260909 -9.617915 25% -0.319611 -0.071712 5.980000 -0.328476 -0.053379 50% 0.015231 -0.049750 0.000914 0.010753 22.350000

0.090329

11.135740

0.076267

15.373170 19656.530000

78.930000

75%

max

0.351466

5.541598

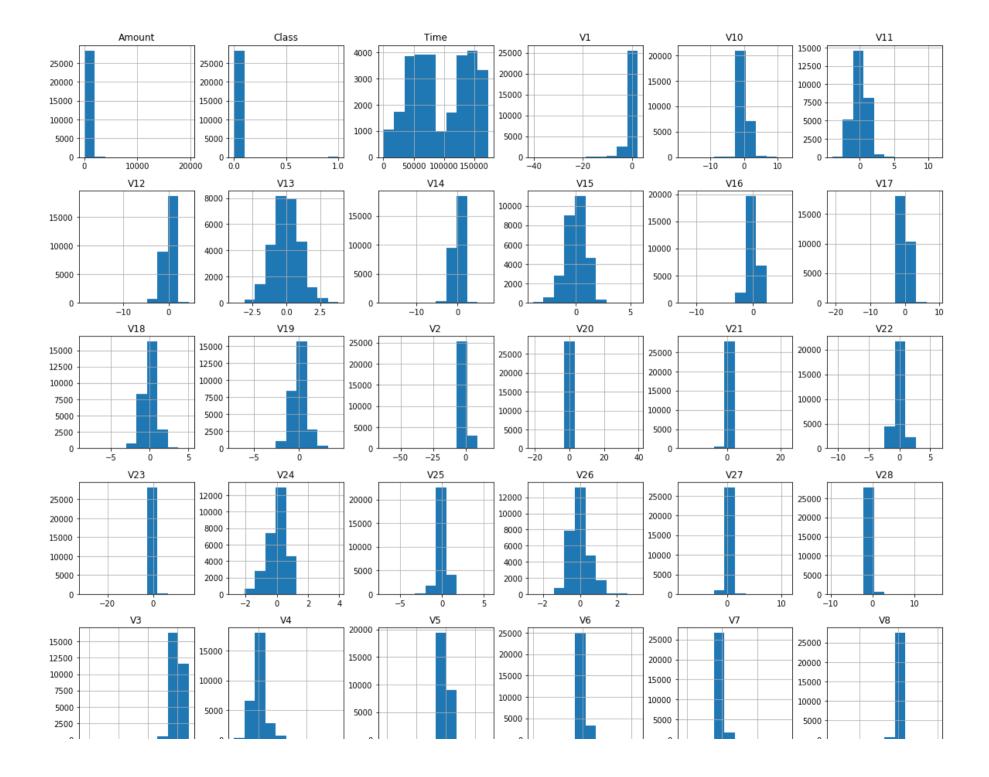
0.253580

3.118588

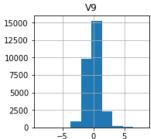
Class count 28481.000000 0.001720 mean std 0.041443 min 0.000000 25% 0.000000 50% 0.000000 75% 0.000000 1.000000 max

[8 rows x 31 columns]

```
In [6]: # Plot histograms of each parameter
    data.hist(figsize = (20, 20))
    plt.show()
```







```
In [7]: # Determine number of fraud cases in dataset

Fraud = data[data['Class'] == 1]
   Valid = data[data['Class'] == 0]

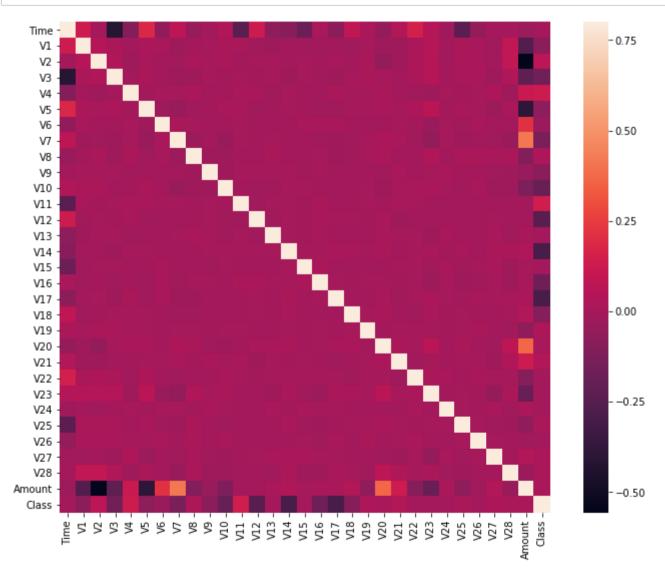
outlier_fraction = len(Fraud)/float(len(Valid))
   print(outlier_fraction)

print('Fraud Cases: {}'.format(len(Fraud)))
   print('Valid Transactions: {}'.format(len(Valid)))
```

0.0017234102419808666

Fraud Cases: 49

Valid Transactions: 28432



```
In [9]: # Get all the columns from the dataFrame
    columns = data.columns.tolist()

# Filter the columns to remove data we do not want
    columns = [c for c in columns if c not in ["Class"]]

# Store the variable we'll be predicting on
    target = "Class"

X = data[columns]
Y = data[target]

# Print shapes
print(X.shape)
print(Y.shape)

(28481, 30)
(28481,)
```

## **The Algorithms**

## **Local Outlier Factor (LOF)**

The anomaly score of each sample is called Local Outlier Factor. It measures the local deviation of density of a given sample with respect to its neighbors. It is local in that the anomaly score depends on how isolated the object is with respect to the surrounding neighborhood.

## **Isolation Forest Algorithm**

The IsolationForest 'isolates' observations by randomly selecting a feature and then randomly selecting a split value between the maximum and minimum values of the selected feature.

Since recursive partitioning can be represented by a tree structure, the number of splittings required to isolate a sample is equivalent to the path length from the root node to the terminating node.

This path length, averaged over a forest of such random trees, is a measure of normality and our decision function.

Random partitioning produces noticeably shorter paths for anomalies. Hence, when a forest of random trees collectively produce shorter path lengths for particular samples, they are highly likely to be anomalies.

```
In [10]: from sklearn.metrics import classification_report, accuracy_score
         from sklearn.ensemble import IsolationForest
                                                                     #anomalv detection
         from sklearn.neighbors import LocalOutlierFactor #anomalv detection
         #localOutlierFactor is an unsupervised outlier detection method
         #It calculates the anomaly score of each sample and we call it the localOutlierFactor
         #It measures the local deviation of density of a given sample wrt it's neighbors
         #It is local and that anomaly score depends on how isolated the object is wrt the surrounding neighborhood
         #It is determined in the same way as Kneighbors method
         #Isolation Forest explicitly identifies anomalies instead of profiling normal data points
         #Isolation Forest, like any tree ensemble method, is built on the basis of decision trees
         #In these trees, partitions are created by first randomly selecting a feature and then
         #selecting a random split value between the minimum and maximum value of the selected feature
         #In principle, outliers are less frequent than regular observations and are different from them in terms of values
         #they lie further away from the regular observations in the feature space
         #define a random state
         state = 1
         #define the outlier detection methods
         #putting into a dictionary of classifiers
         classifiers = {
              "Isolation Forest": IsolationForest(
                                              \max \text{ samples} = \text{len}(X),
                                              contamination = outlier fraction,
                                              random state = state),
              "Local Outlier Factor": LocalOutlierFactor(
                                             n_neighbors = 20,
```

contamination = outlier\_fraction)

```
In [14]: # Fit the model
         #plt.figure(figsize=(9, 7))
         n_outliers = len(Fraud)
         for i, (clf name, clf) in enumerate(classifiers.items()):
             # fit the data and tag outliers
             if clf name == "Local Outlier Factor":
                 y pred = clf.fit predict(X)
                 scores_pred = clf.negative_outlier_factor_
             else:
                 clf.fit(X)
                 scores pred = clf.decision function(X)
                 y pred = clf.predict(X)
             # Reshape the prediction values to 0 for valid, 1 for fraud.
             y pred[y pred == 1] = 0
             y pred[y pred == -1] = 1
             n errors = (y pred != Y).sum()
             # Run classification metrics
             print('{}: {}'.format(clf name, n errors))
             print(accuracy_score(Y, y_pred))
             print(classification report(Y, y pred))
```

C:\ProgramData\Anaconda3\lib\site-packages\sklearn\ensemble\iforest.py:247: FutureWarning: behaviour="old" is depreca ted and will be removed in version 0.22. Please use behaviour="new", which makes the decision\_function change to match other anomaly detection algorithm API.

FutureWarning)

C:\ProgramData\Anaconda3\lib\site-packages\sklearn\ensemble\iforest.py:415: DeprecationWarning: threshold\_ attribute is deprecated in 0.20 and will be removed in 0.22.

" be removed in 0.22.", DeprecationWarning)

Isolation Forest: 71

weighted avg

0.99750711000	316			
	precision	recall	f1-score	support
0	1.00	1.00	1.00	28432
1	0.28	0.29	0.28	49
accuracy			1.00	28481
macro avg	0.64	0.64	0.64	28481
weighted avg	1.00	1.00	1.00	28481
Local Outlier 0.99659422070				
	precision	recall	f1-score	support
0	1.00	1.00	1.00	28432
1	0.02	0.02	0.02	49
accuracy			1.00	28481
macro avg	0.51	0.51	0.51	28481

1.00

1.00

28481

1.00