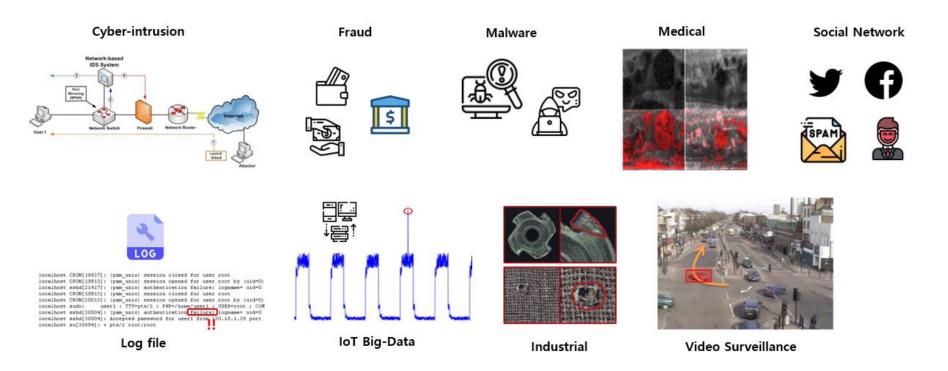
Credit Card Fraud Detection

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Examples of anomaly detection



Anomaly Detection

Supervised Anomaly Detection:

Supervised anomaly detection techniques require a data set that has been labeled as "normal" and "abnormal" and involves training a classifier. However, this approach is rarely used in anomaly detection due to the general unavailability of labelled data and the inherent unbalanced nature of the classes.

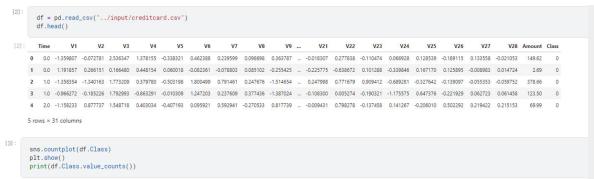
Semi-supervised (One-Class) Anomaly Detection

Semi-supervised anomaly detection techniques assume that some portion of the data is labelled. This may be any combination of the normal or anomalous data, but more often than not the techniques construct a model representing normal behavior from a given *normal* training data set, and then test the likelihood of a test instance to be generated by the model.

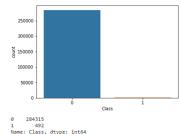
Unsupervised Anomaly Detection

Unsupervised anomaly detection techniques assume the data is unlabelled and are by far the most commonly used due to their wider and relevant application.

```
import pandas as pd
 import numpy as np
 import seaborn as sns
 import matplotlib.pyplot as plt
 import matplotlib.gridspec as gridspec
 #for data preprocessing
 from sklearn.decomposition import PCA
  #for modeling
 from sklearn.neighbors import LocalOutlierFactor
 from sklearn.ensemble import IsolationForest
 #filter warnings
 import warnings
 warnings.filterwarnings("ignore")
 import os
 print(os.listdir("../input"))
 # Any results you write to the current directory are saved as output.
['creditcard.csv']
```

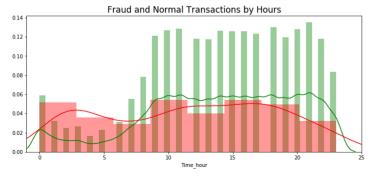


Data that has been converted to V1 to V28 by PCA was used due to confidentiality issues.



```
timedelta = pd.to_timedelta(df['Time'], unit='s')
df['Time_hour'] = (timedelta.dt.components.hours).astype(int)

plt.figure(figsize=(12,5))
sns.distplot(df[df['Class'] == 0]["Time_hour"], color='g')
sns.distplot(df[df['Class'] == 1]["Time_hour"], color='r')
plt.title('Fraud and Normal Transactions by Hours', fontsize=17)
plt.xlim([-1,25])
plt.show()
```



PCA.fit(): found the principal

PCA.transform(): Transform data into new principal components

```
[6]
       columns = df.drop('Class', axis=1).columns
                                                                                                                                                             0.25 -
       grid = gridspec.GridSpec(6, 5)
                                                                                                                                                            015
       plt.figure(figsize=(20,10*2))
                                                                                                                                                                                                              -1010
       for n, col in enumerate(df[columns]):
            ax = plt.subplot(grid[n])
            sns.distplot(df[df.Class==1][col], bins = 50, color='g')
            sns.distplot(df[df.Class==0][col], bins = 50, color='r')
            ax.set_ylabel('Density')
            ax.set_title(str(col))
                                                                                                                     0.0 -20 0 20 40 00
                                                                                                                                                                  -60 -40 VI320 0 22
            ax.set_xlabel('')
       plt.show()
                                                                                                                                                                -50 -25 Via 21 50 75
                                                                                                                                                  -1P<sub>17</sub>
                                                                                                                    E 0.75
                                                                                                                     1.00
                                                                                                                    £ 0.75 -
```

```
[7]:
       def ztest(feature):
                                                                                                                             z-test:
            mean = normal[feature].mean()
            std = fraud[feature].std()
            zScore = (fraud[feature].mean() - mean) / (std/np.sqrt(sample_size))
                                                                                                                            The z-test is an analytical technique that tests a
                                                                                                                            hypothesis by comparing the means of two groups.
            return zScore
      columns= df.drop('Class', axis=1).columns
      normal= df[df.Class==0]
                                                                                                                            Valid transactions as our population
      fraud= df[df.Class==1]
      sample_size=len(fraud)
                                                                                                                            Fraud transactions as sample
      significant_features=[]
      critical_value=2.58
                                                                                                                            Two tailed Z-test
      for i in columns:
                                                                                                                            Level of significance 0.01
         z vavlue=ztest(i)
                                                                                                                            Corresponding critical value is 2.58
         if( abs(z_vavlue) >= critical_value):
             print(i," is statistically significant") #Reject Null hypothesis. i.e. H0
             significant_features.append(i)
     V1 is statistically significant
                                                                                                                            Hypothesis:
     V2 is statistically significant
     V3 is statistically significant
     V4 is statistically significant
     V5 is statistically significant
                                                                                                                            H0: There is no difference (insignificant)
     V6 is statistically significant
    V7 is statistically significant
     V9 is statistically significant
                                                                                                                            H1: There is a difference (significant)
     V10 is statistically significant
     V11 is statistically significant
     V12 is statistically significant
     V14 is statistically significant
     V16 is statistically significant
    V17 is statistically significant
    V18 is statistically significant
     V19 is statistically significant
     V20 is statistically significant
                                                                                                                           Zscore=(x^-\mu)/S.E
     V21 is statistically significant
     V24 is statistically significant
     V27 is statistically significant
     V28 is statistically significant
     V29 is statistically significant
    V30 is statistically significant
```

```
significant_features.append('Class')
      df= df[significant_features]
      inliers = df[df.Class==0]
      ins = inliers.drop(['Class'], axis=1)
      outliers = df[df.Class==1]
      outs = outliers.drop(['Class'], axis=1)
      ins.shape, outs.shape
[9]: ((284315, 23), (492, 23))
      def normal_accuracy(values):
         tp=list(values).count(1)
         total=values.shape[0]
         accuracy=np.round(tp/total,4)
         return accuracy
      def fraud_accuracy(values):
         tn=list(values).count(-1)
         total=values.shape[0]
         accuracy=np.round(tn/total,4)
          return accuracy
```

Modeling

```
IForest
 state= 42
  ISF = IsolationForest(random_state=state)
  ISF.fit(ins)
                                                                                                       Scores
                                                                                                                                                                                                    ITree
  normal_isf = ISF.predict(ins)
 fraud_isf = ISF.predict(outs)
                                                                                         Outlier
                                                                                 Normal uncommon
  in_accuracy_isf=normal_accuracy(normal_isf)
                                                                                      samples
  out_accuracy_isf=fraud_accuracy(fraud_isf)
  print("Accuracy in Detecting Normal Cases:", in_accuracy_isf)
  print("Accuracy in Detecting Fraud Cases:", out_accuracy_isf)
                                                                                  Normal common
Accuracy in Detecting Normal Cases: 0.9
                                                                                      samples
Accuracy in Detecting Fraud Cases: 0.9004
```

IsolationForest

It is mainly used to detect outliers in the current data. As the name suggests, it is implemented based on a tree, and it is implemented by splitting the data at random and isolating all observations.

In particular, it has the advantage of being able to operate efficiently on data with many variables.

Modeling

```
LOF = LocalOutlierFactor(novelty=True)
LOF.fit(ins)

normal_lof = LOF.predict(ins)
fraud_lof = LOF.predict(outs)

in_accuracy_lof=normal_accuracy(normal_lof)
out_accuracy_lofsfraud_accuracy(fraud_lof)
print("Accuracy in Detecting Normal Cases:", in_accuracy_lof)
print("Accuracy in Detecting Normal Cases:", out_accuracy_lof)

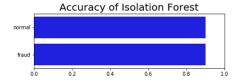
Accuracy in Detecting Normal Cases: 0.9171
Accuracy in Detecting Fraud Cases: 0.5142
```

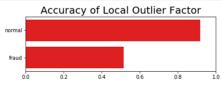
LocalOutlierFactor

LOF indicates how far each observation is within the data (outliers). The most important characteristic of LOF is not to consider all data as a whole, but rather to use the data surrounding that observation to identify the extent of an outlier from a local point of view.

Result

```
[13]:
       fig, (ax1,ax2)= plt.subplots(1,2, figsize=[15,2])
       ax1.set_title("Accuracy of Isolation Forest",fontsize=20)
       sns.barplot(x=[in_accuracy_isf,out_accuracy_isf],
                   y=['normal', 'fraud'],
                   label="classifiers",
                   color="b",
                   ax=ax1)
       ax1.set(xlim=(0,1))
       ax2.set_title("Accuracy of Local Outlier Factor", fontsize=20)
       sns.barplot(x=[in_accuracy_lof,out_accuracy_lof],
                   y=['normal', 'fraud'],
                  label="classifiers",
                   color="r",
                   ax=ax2)
       ax2.set(xlim=(0,1))
       plt.show()
```





Conclusion

Both, Isolation Forest and Local Outlier Factor performed same in predicting Normal cases but Isolation Forest performed far better in detecting Fraud cases.

Reference

https://www.kaggle.com/code/sabanasimbutt/anomaly-detection-using-unsupervised-techniques

"Deep Learning for Anomaly Detection: A Survey," 2019 arXiv